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# NAVAL POSTGRADUATE SCHOOL

## Monterey, California



CONTRACTOR REPORT

GAS DYNAMICS OF LASER EXHAUST EXTERNAL TO SPACECRAFT

Shaul Abramovich

November 1985

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The work reported herein was carried out for the Naval Postgraduate School by Dr. Shaul Abramovich under Contract Number N62271-83-M-1939. The work presented in this report is in support of DARPA project. The work is based on general publications and theory and provides specific means for solving the exhaust flow from a spacecraft laser. Computer programs have been developed to calculate the flowfield in the continuum region as well as in the molecular region. It provides the means to calculate the flux of the exhausted gas towards the walls of the spacecraft. The project on LASER EXHAUST is funded by Defense Advanced Research Projects Agency and is under the cognizance of Distinguished Professor A. E. Fuhs.

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## 20. Abstract (continued)

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The axisymmetric Monte Carlo Simulation begins outside the nozzle, where the breakdown parameter has a value of approximately 0.05. The actual shape of this breakdown limit is a closed lobe surface which starts at the nozzle lips, approximately follows a specific Mach wave in the Prandtl-Meyer fan and closes back to the axis far downstream. Because our interest is limited to the close vicinity of the corners, there the shape of this limit may be approximated to a straight line (for an axisymmetric flow making a cone).

For the simulation purposes, the domain between the breakdown limit and the wall is divided into sectors, each sector divided into radial regions and each region into simulation cells. Each cell is filled with a number of simulated molecules relative to the cell volume and local number density.

The simulation is performed for each region separately and contains:

- \* molecular motions
- \* generation of new molecules to simulate input flows
- \* deactivation of molecules to simulate output flows
- \* molecular collisions

Because there is no apriori information about the flow interaction between different regions, the whole simulation is done on an iterative basis. A first run will supply the output flux from each region. These results are used as input data for consecutive runs.

The program runs as far as the collision frequency is still high and the mean free path is low compared with the size of the cells. Beyond this limit the flow may be regarded as collisionless, and the flux towards the solid wall may be computed directly.

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## ABSTRACT

Some procedures have been developed to analyze the flowfield of highly underexpanded axisymmetric ring jets operated at high altitudes. The Method of Characteristics (MOC) was used to compute the Prandtl-Meyer expansion fan and the flow parameters in that region. The MOC may also be used to obtain some indications about the repetitive expansion — compression behavior of the jet as well as the divergent shape of the expansion part downstream, when the ambient pressure goes below certain limits.

The continuum methods may be used as far as the limit where translational equilibrium ceases to exist. The breakdown of the continuum theory may be evaluated using the experimental breakdown parameter as proposed by G. A. Bird. Beyond this limit, the molecular Direct-Simulation-Monte-Carlo method (DSMC) is applied.

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For the simulation purposes, the domain between the breakdown surface and the wall is divided into sectors, each sector divided into radial regions and each region into simulation cells. Each cell is filled with a number of simulated molecules relative to the cell volume and local number density.

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- \* molecular motions
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The program runs as far as the collision frequency is still high and the mean free path is low compared with the size of the cells. Beyond this limit the flow may be regarded as collisionless, and the flux towards the solid wall may be computed directly.



## I. INTRODUCTION

Gas jets released from spacecrafts and external flows about vehicles at high altitudes have a renewed interest in particular with regard to two main aspects:

- a. contamination of the spacecraft walls
- b. optical disturbances caused by the plume.

A spacecraft gas dynamics laser releasing a large quantity of gas is highly affected by these two factors and the interest in analyzing them and being able to control them have a unique importance in further development.

A spacecraft laser is assumed to have a long cylindrical shape with the optical power output devices installed at one of its bases as shown in Figure 1.

The output gas is released through a ring nozzle, undergoes a fast three dimensional axisymmetric expansion, and forms a plume covering the whole meridian plane of the vehicle. It widens to large angles of expansion so that it may intersect the laser beam. Back scattered molecules may return to the wall of the spacecraft causing contamination and degradation of surfaces and vehicle parts.



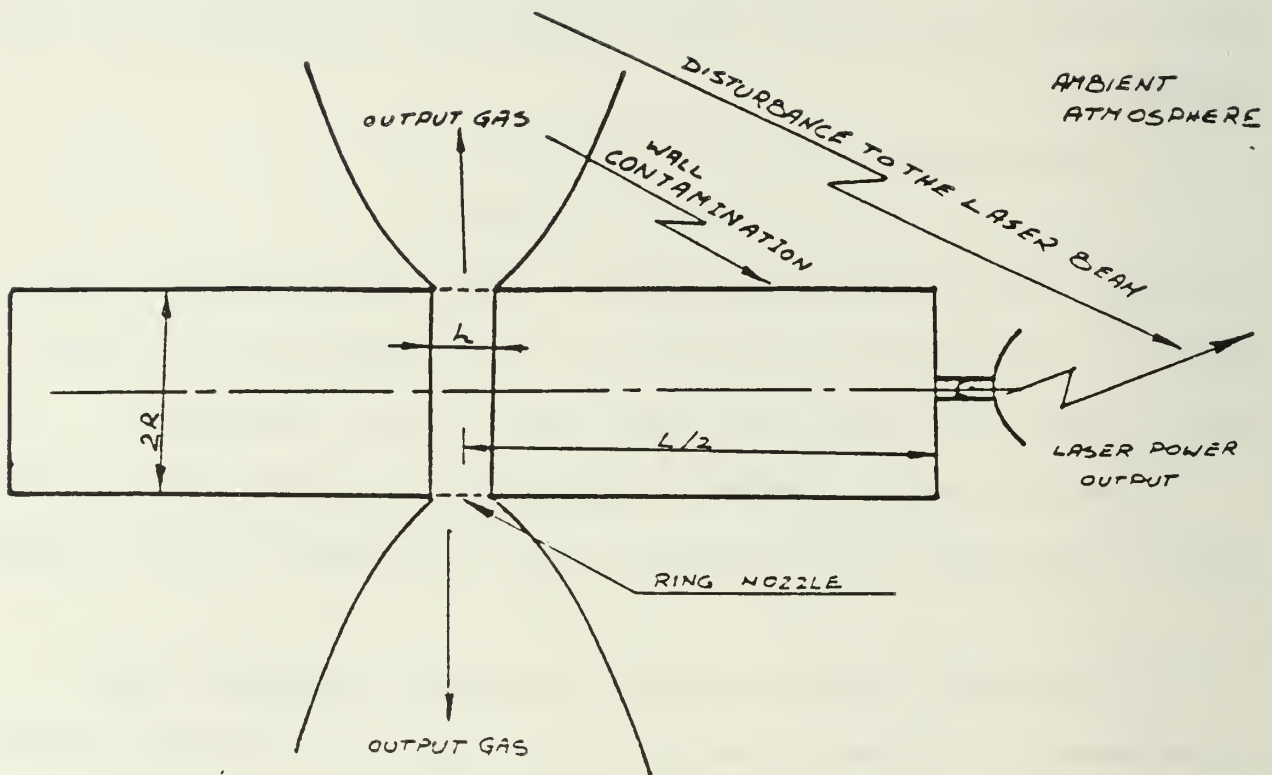


Figure 1. The Spacecraft Laser.

$M_0$  - Mach number at the exit surface  $\sim 4$  the jet gas is composed of two species

heavy molecules  $M_{G1} = 19$

light molecules  $M_{G2} = 4$

Altitudes between 200 to 1000 km.

Continuum flow theory may be used to solve the flowfield and flow parameters as far as there is translational equilibrium, it means that intermolecular interactions are fast enough to maintain expansion rates. Wherever these interactions are too slow, the continuum flow becomes invalid and the molecular flow theory should be employed.

The solution for the continuum regime is computed here by means of the Method of Characteristics (MOC) [1,2,3]. The limit where continuum breakdown occurs was estimated by the experimental breakdown parameter as proposed by G. A. Bird [4]. Beyond this limit, it is proposed to compute the molecular flow by means of the Direct Simulation Monte Carlo technique as described in detail by Bird [4].

For moderate and low pressure ratios (static pressure at the nozzle exit to the ambient pressure), an underexpanded jet exhibits a repetitive expansion - compression behavior with a geometry depending on the initial Mach angle, on the Prandtl-Meyer fan angle, and on the gas specific heats ratio. For lower ambient pressure which occurs at higher altitudes, the first compression region is pushed out to the envelope of the jet forming the barrel shock. If the ambient pressure is low enough, this compression region may disappear due to the molecular behavior of the flow [6,11].

The breakdown of the continuum theory occurs in a region where the gas density and pressure are high compared with the density and pressure in the ambient gas. At high altitudes the ratios between these parameters may reach  $10^6$  or more. Considering this range of variations, computational validity dictates the use of the Direct Simulation Monte Carlo method. In the higher density range the jet will be considered as consisting of two species of gas, their molecular model will be "the hard sphere molecule" model and

ambient gas is not allowed to protrude. In the lower density region the flow will be regarded as collisionless.

In the following chapters we bring the detailed description of the computer programs which solve the different parts of the flowfield.

## II. THE CONTINUUM REGIME

### A. THE TWO DIMENSIONAL ISENTROPIC UNDEREXPANDED JET

The results brought here are based on the supersonic steady isentropic flow theory as described in literature (see for example, Shapiro [1], Liepman and Roshko [2], and Owczarek [3]).

The ranges of parameters of a jet emerging from a gas dynamics spacecraft laser are:

a. The Mach number at the exit surface  $M_0 \approx 4$ . The static pressure at the exit plane  $P_0 \approx 136\text{pa}$ . Ambient pressure ( $P_{\text{amb}}$ ), temperature ( $T_{\text{amb}}$ ) and other thermodynamic properties of ambient gas depend on the altitude as shown in Table 1. The jet gas may consist of DF, HF, Helium and other species. In the programs we limit the composition to two species: Air and He. (The program allows changes in the composition and types of gas.)

b. The pressure ratio  $\frac{P_0}{P_{\text{amb}}}$  for the minimum required altitude (200 km) assures that the jet is highly underexpanded (we show later the influence of this ratio on the shape of the jet).

The following thermodynamic relations are valid as long as the compressible flow is isentropic

$$T_T = T(1 + \frac{\gamma-1}{2} M^2) \quad (1)$$

$$P_T = P(1 + \frac{\gamma-1}{2} M^2)^{\frac{\gamma}{\gamma-1}} \quad (2)$$

$$\rho_T = \rho(1 - \frac{\gamma-1}{2} M^2)^{\frac{1}{\gamma-1}} \quad (3)$$

TABLE 1

ATMOSPHERIC DATA  
(Abstracted from U.S. Standard Atmosphere 1976)

Altitude km	Pressure m bar	number density m <sup>-3</sup>	particle speed m/sec	collision frequency sec <sup>-1</sup>	mean free path m	molecular weight kg/kmol	density kg/m <sup>3</sup>	temperature °k
100	3.2011 -4	1.189 +19	381.4	2.68 +3	1.42 -1	28.4	5.604 -7	195.08
200	8.4736 -7	7.182 +15	921.6	3.9	2.4 +2	21.3	2.541 -10	854.56
300	8.7704 -8	6.509 +14	1079.7	4.2 -1	2.6 +3	17.73	1.916 -11	976.01
400	1.4518 -8	1.056 +14	1148.5	7.2 -2	1.6 +4	15.98	2.803 -12	995.83
500	3.0236 -9	2.192 +13	1215.0	1.6 -2	7.7 +4	14.33	5.215 -13	999.24
600	8.2130 -10	5.950 +12	1356.4	4.8 -3	2.8 +5	11.51	1.137 -13	999.85
700	3.1908 -10	2.311 +12	1627.0	2.2 -3	7.3 +5	8.00	3.070 -14	999.97
800	1.7036 -10	1.234 +12	1954.3	1.4 -3	1.4 +6	5.54	1.136 -14	999.99
900	1.0873 -10	7.876 +11	2192.6	1.0 -3	2.1 +6	4.40	5.759 -15	1000.00
1000	7.5138 -11	5.442 +11	2318.1	7.5 -4	3.1 +6	3.94	3.561 -15	1000.00



where  $T_T$ ,  $P_T$ , and  $\rho_T$  are the total temperature, pressure and density (constant for isentropic field).  $T$ ,  $P$ , and  $\rho$  are local temperature, pressure and density  $\gamma$  is the specific heat ratio of the gas (considered here as constant),  $M$  is the local Mach number.

The partial differential equation of motion for supersonic 2-D irrotational and isentropic flow is a hyperbolic equation having solutions obtained from invariants along characteristic lines. Physical interpretation of these lines are the compression or expansion waves which are oriented at Mach angles relative to the streamlines.

Once the directions of the characteristics (waves) are determined everywhere in the field, all other parameters may be calculated.

#### B. THE TWO DIMENSIONAL PLANAR JET

The compressible supersonic jet flow is characterized by two families of characteristics (pressure waves) starting at each corner of the nozzle lips. Each of these families of waves forms a Prandtl-Meyer fan. The streamlines crossing the characteristic waves bend outwards resulting in an increase in the flow area. The angle  $\mu$  between the streamline and the pressure wave is a function of the local Mach number as

$$\mu = \arcsin (1/M) \quad (4)$$

The symbol  $\mu$  is the Mach angle.

Using the isentropic relations, we can find the relation between the turning angle ( $\theta$ ) and the local Mach number ( $M$ ) as

$$d\theta = - \frac{(M^2 - 1)}{M(1 + \frac{\gamma-1}{2} M^2)} dM \quad (5)$$

Integration between the conditions  $M=1$  and  $M$  gives the total turning angle starting at the throat (where  $M=1$ ) up to a point with given  $M$ . The result gives the Prandtl-Meyer function (angle) as

$$v(M) = \frac{\gamma+1}{\gamma-1} \arctg \frac{\gamma-1}{\gamma+1} (M^2-1) - \arctg (M^2-1) \quad (6)$$

In the close vicinity of the nozzle lips where the two families of characteristics do not intersect with each other there is a "simple region" of expansion. There the flow parameters are defined by  $v$  and  $\theta$  of each characteristic line. Further downstream the waves intersect each other. In this part of the flow parameters are defined by the two intersecting characteristics. A singularity occurs when the initial Mach number is unity and a special treatment is required to start the calculations at that point (this special treatment has not been brought here). A particular importance of the Prandtl-Meyer function is when analyzing the two dimensional flow using the hodograph plane.

#### C. THE HODOGRAPH PLANE FOR A TWO DIMENSIONAL JET

The hodograph plane is a representation of the flow parameters in the velocity plane. Figure 2 shows a hodograph plane calculated for a simple gas ( $\gamma=1.4$ ). The circles represent constant Mach numbers and constant

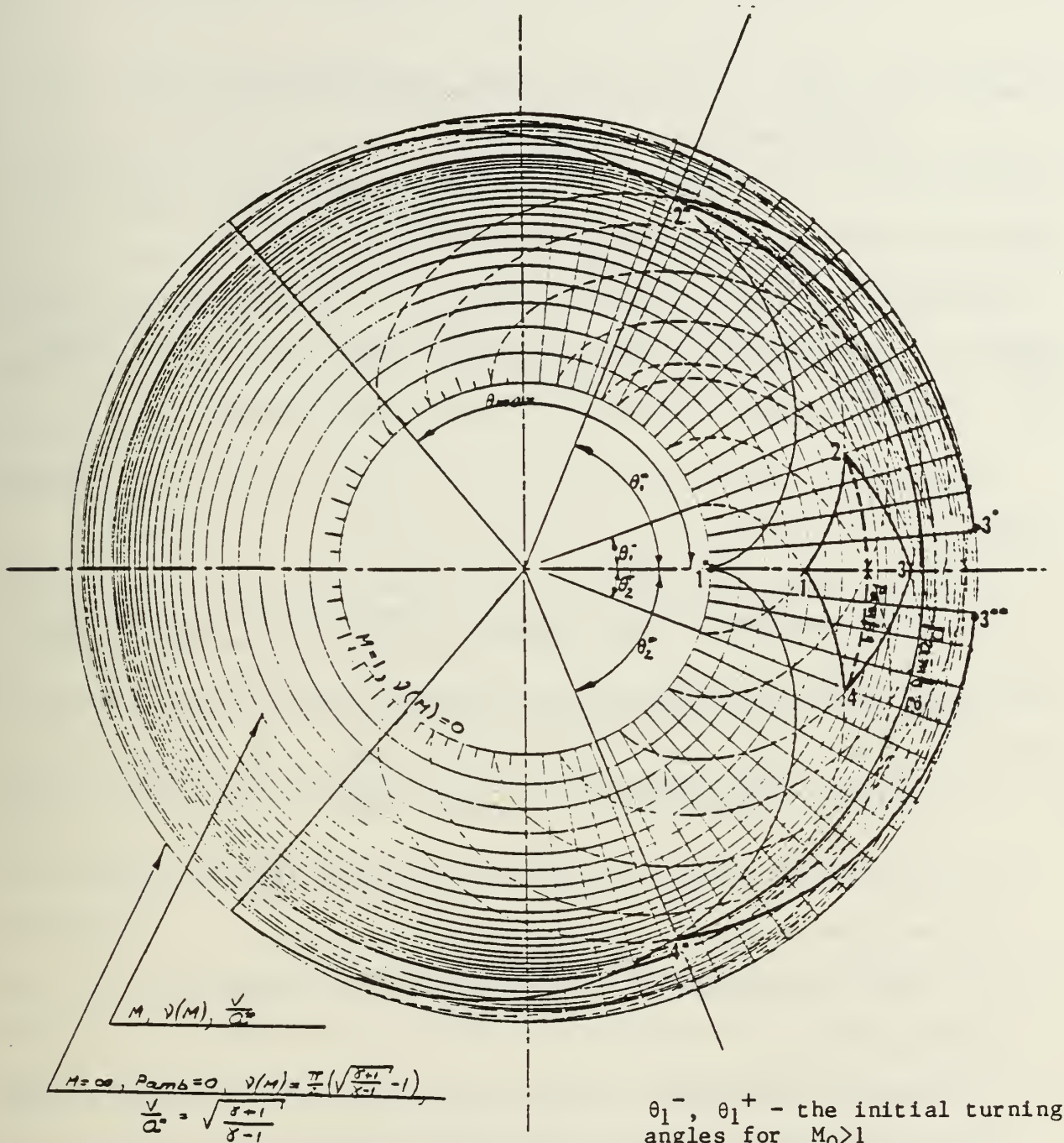


Figure 2. The Hodograph Plane.

1-2-3-4-1 defines a jet with  $M_0 > 1$  expanding into  $P_{amb1} < P_0$ , resulting simple repetitive expansion-compression.  
 1\*-2\*-3\*-3\*\*-4\*-1\* defines a jet with  $M_0 = 1$  expanding into  $P_{amb2} \ll P_0$ , resulting in a highly underexpanded jet.

pressure ratios  $(\frac{P}{P_T})$  . The epicycloids are the two families of characteristics. The angles  $\theta$  are the turning angles due to expansion or compression (which are both present in supersonic jets).

To define an isentropic supersonic jet on the hodograph plane it is necessary to define the Mach number at the exit plane, the pressure ratios  $(\frac{P}{P_T})$  and  $(\frac{P_{amb}}{P_T})$  , and the specific heat ratio ( $\gamma$ ) of the gas.

Investigating the shapes of the jet as a function of ranges of parameters we may get:

- a. simple underexpanded jets for which  $\frac{P_{amb}}{P_T}$  is high enough so that the two families of characteristics intersect each other.
- b. a critical underexpanded jet for which  $\frac{P_{amb}}{P_T}$  is low enough so that the intersection between the outer characteristic lines of the two families occur on the outer hodograph circle.
- c. highly underexpanded jets for which a part of characteristics do not intersect at all.
- d. expansion into complete vacuum so that there are no reflections from the jet boundaries and therefore the compression region disappears.

## D. THE SHAPES OF TWO DIMENSIONAL JETS

The following paragraphs further detail the different shapes.

### 1. Simple Underexpanded Jets

Figure 3 shows the physical plane and the hodograph plane of a simple underexpanded jet. If  $M_0 > 1$ , an initial turn of the flow  $\theta_0$  is made within the nozzle. An additional turn of  $\theta$  is due to the underexpansion.  $\theta$  is found by the intersection of characteristics (1-2) with the circle defined by  $P_{amb}/P_T$ .

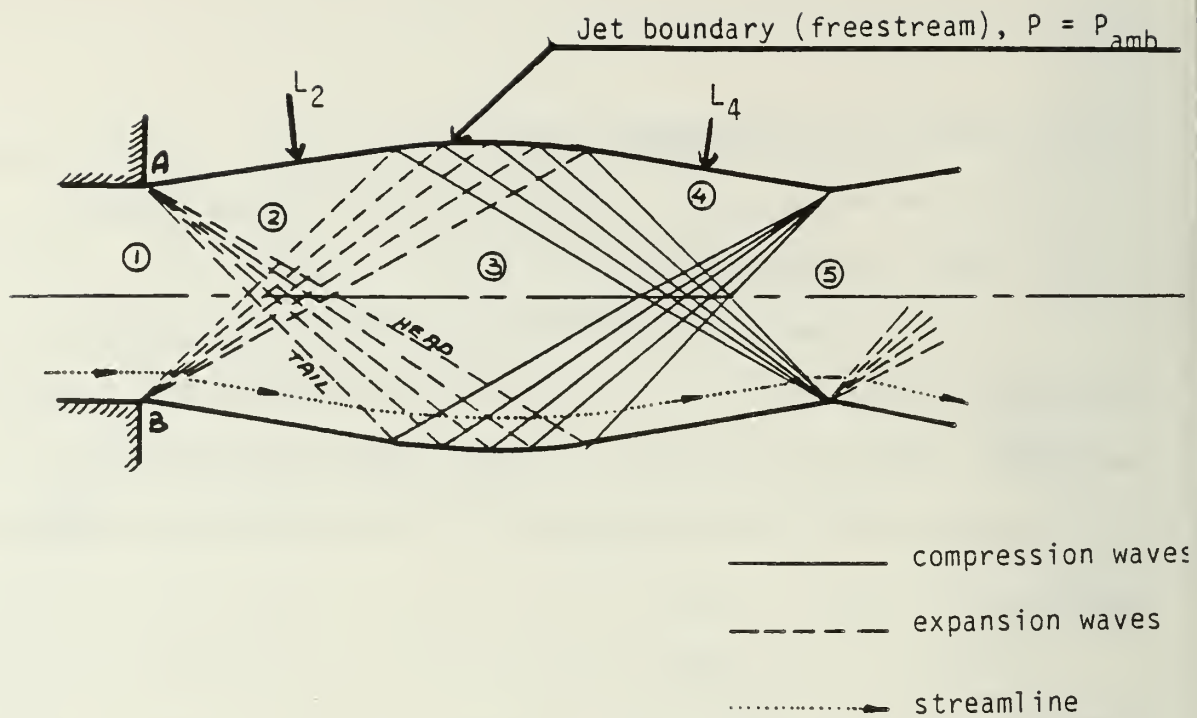
When  $M_0 > 1$ , the characteristic line 1-2 is described in the physical plane by a region in which only one family of expansion waves are present (simple region, see transverse line between 1 to 2 in the physical plane). A different family of expansion waves forms a second simple region when moving between 2 to 3. At a larger distance from the exit plane, reflected waves from the free streamline (jet boundary) cause compression. An ideal representation of such a jet is a repetitive pattern of expansion and compression.

For  $M_0 = 1$ , the tail waves of both families are perpendicular to the flow, thus, both lie on line AB. That means that if  $M_0 = 1$  there is no simple region near the exit plane of the jet. The characteristic line 1-2 on the hodograph plane becomes a single point A (or B) when located in the physical plane.

### 2. Critical Underexpanded Jets

We define a critical underexpanded jet when point (3) (see Figure 4) lies on the limiting circle  $M_\infty$  or  $P/P_\infty = 0$ . This means that there is a core within the jet where the pressure approaches zero and Mach approaches infinity. This core is theoretically bounded at its upstream side by expansion waves and downstream by compression waves.





- ① uniform axial flow ( $P=P_0$ )
- ② uniform flow parallel to  $L_2$  ( $P=P_{amb}$ )
- ③ uniform axial flow (lowest pressure- $P_{min}$ )
- ④ uniform flow parallel to  $L_4$  ( $P=P_{amb}$ )
- ⑤ uniform axial flow ( $P=P_0$ )

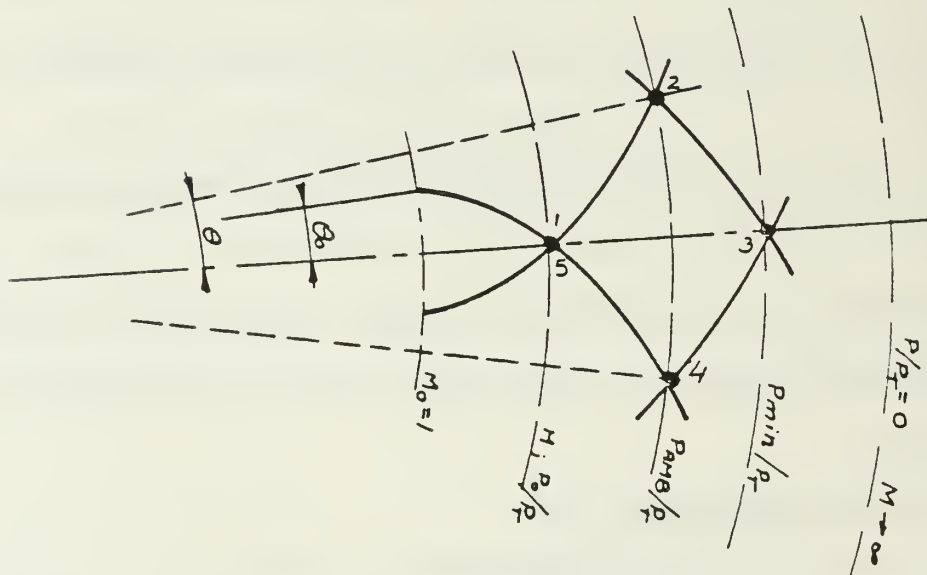


Figure 3. Flow at exit of a simple underexpanded jet.

- (a) physical Plane
- (b) Hodograph Plane

Using the theoretical expressions for isentropic ideal flow one may derive the values of  $\frac{P_{amb}}{P_T}$  or  $\frac{P_{amb}}{P_o}$  as function of  $M_o$  which causes a jet to be critical underexpanded.

Figure 5 shows results of  $\frac{P_{amb}}{P_o}$  as functions of  $M_o$  for gases with different specific heat ratio.

### 3. Highly Underexpanded Jets

When  $P_{amb}$  is lower than the critical values as shown in figure 4, point 3 does not exist (there is no intersection between lines 2-3 2'-3'). This means that the repetitive reversible expansion/compression shapes ceases to exist. The envelope of the jet starts at an angle defined by  $\theta$  at the nozzle exit plane and approaches an asymptotic angle defined by  $\theta_{lim}$  (see Figure 6).

In this case we get two (symmetric) groups of characteristics  $C_1-C_2$  and  $C'_1-C'_2$  (Figure 6) with no intersection between them.  $C_1$  and  $C'_1$  define the inner limit for reflected characteristics,  $C_2$  and  $C'_2$  define the outer limit. As the rest of reflected waves lie between  $C_2$  or  $C'_2$  and the jet boundary, we may conclude that a compression region may exist only in a layer along the jet boundary.

Because of irreversible effects such as shear stresses, heat transfer due to high temperature gradients, or condensation effects in real gases, the compression layer may be interpreted as the "barrel shock".

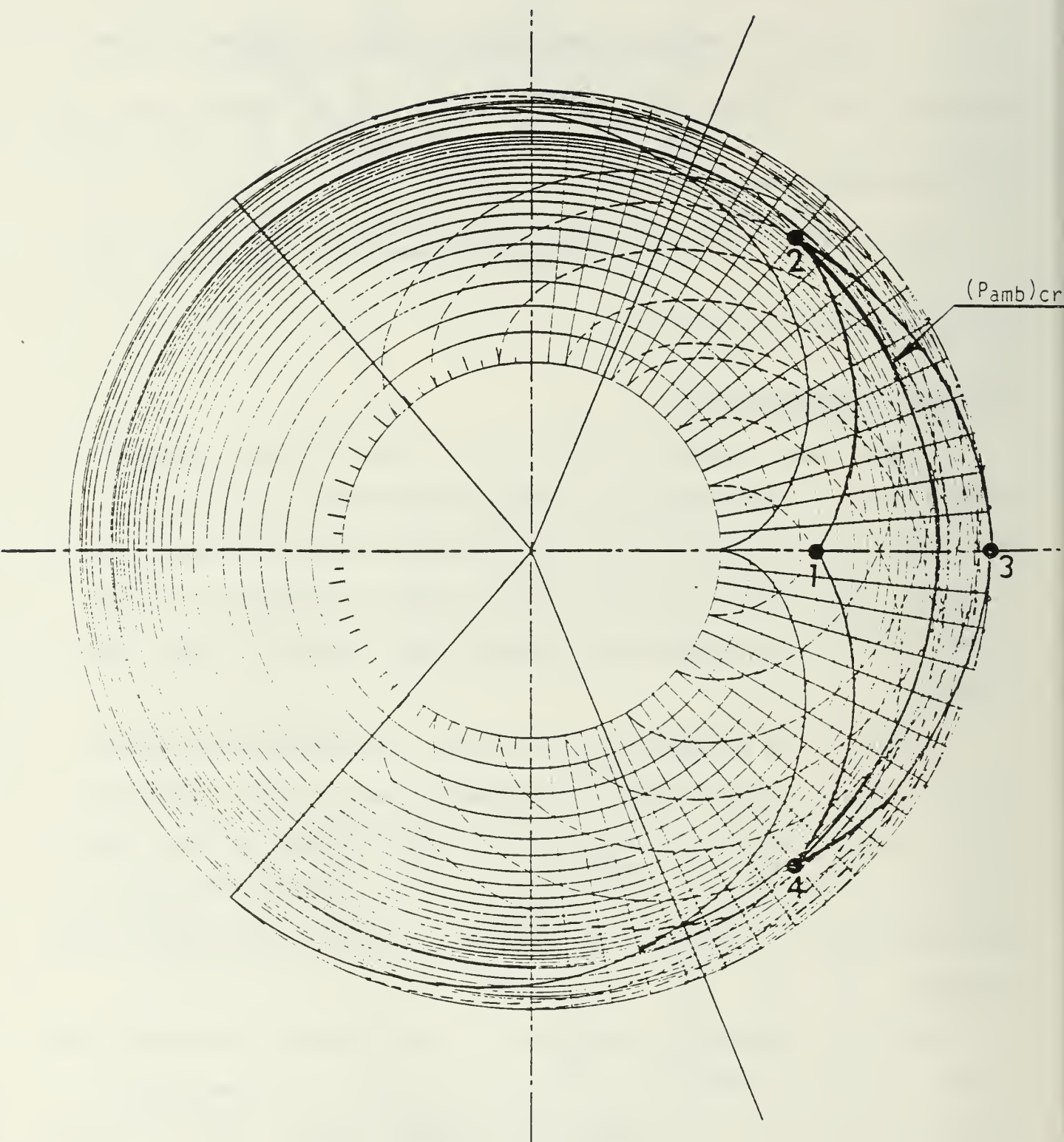


Figure 4. The Hodograph Plane for a Critical Underexpanded Jet.

(1-2-3-4-1), ( $\gamma=1.4$ )

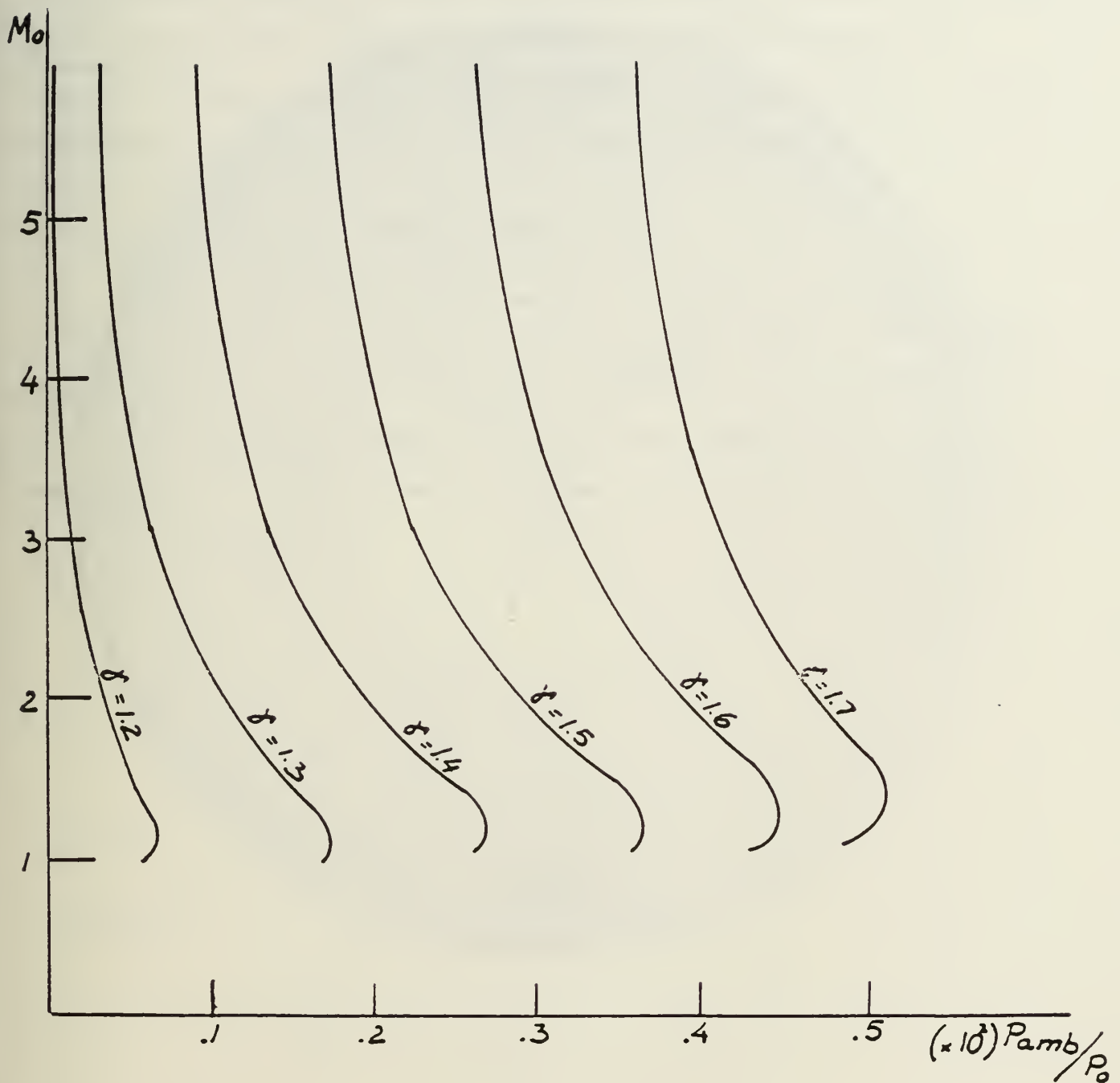


Figure 5. Dependence of critical values of  $P_{amb}/P_0$  on the Mach number at the exit plane ( $M_0$ ) for different values of specific heat ratio ( $\gamma$ ).



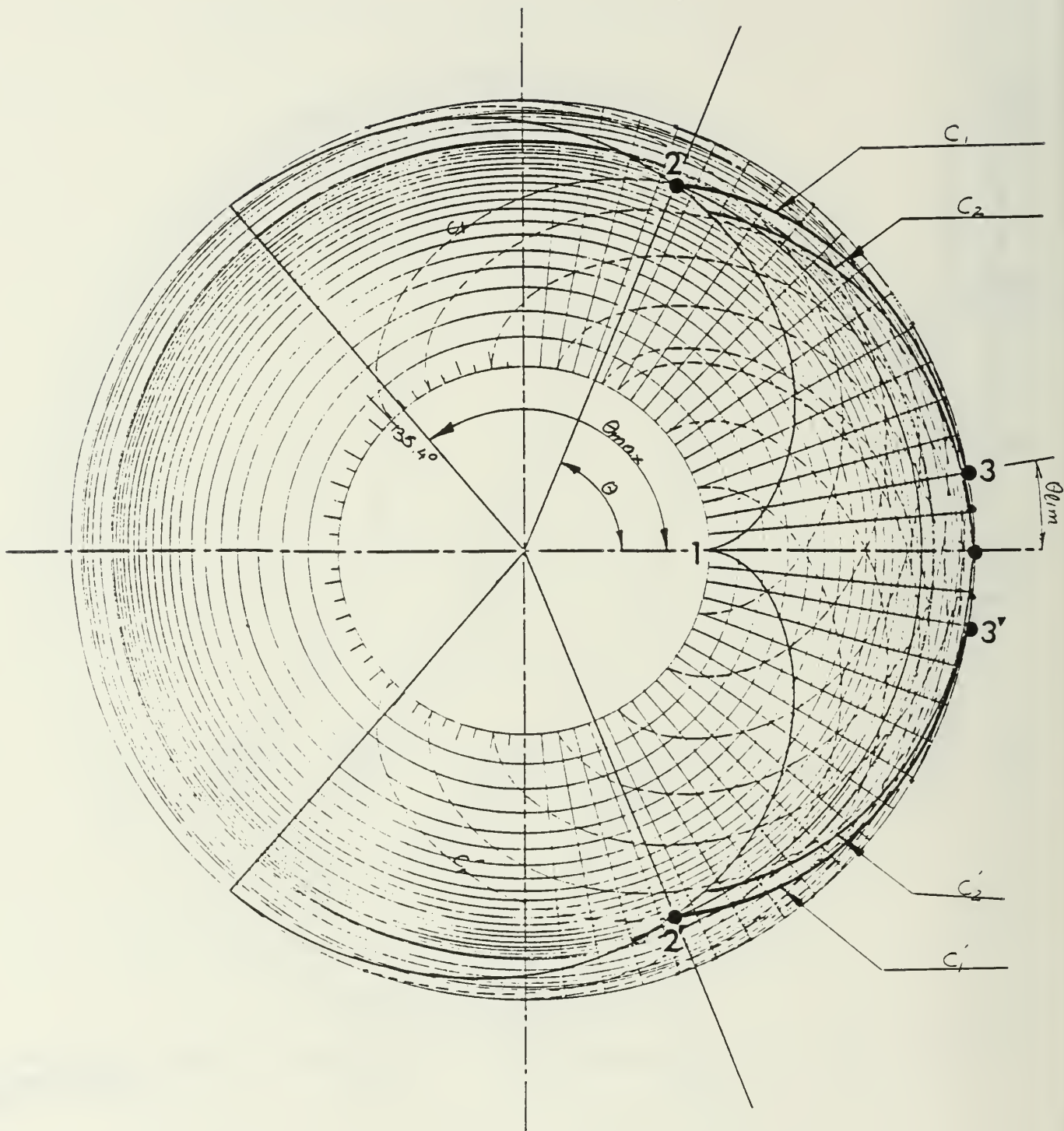


Figure 6. The Hodograph Plane for a Highly Underexpanded Jet

( $\gamma = 1.4$ )

$\theta_{\max}$  - maximum turning angle

$\theta$  - total turning angle in the specific jet  $\theta > \theta_{\max}/2$

$\theta_{\lim}$  - limiting angle of compression region.



Figure (7) shows the hodograph plane for an air jet ( $\gamma=1.4$ ) expanding into an ambient pressure 105 times lower than the static pressure at the exit plane. Figure (8) is a schematic description of the jet (the data for this jet is given in Table 2). The shape shown in Figure (8) may be compared with the barrel shock photograph in page 208 Reference [3].

#### 4. EXPANSION INTO A COMPLETE VACUUM - $P_{amb}=0$

This is an extreme situation in which the maximum turn angle of streamlines occur near the nozzle exit plane. The theoretical free stream is defined only by the theoretical turning angles. All streamlines in the flow expanded monotonically towards  $P \approx 0$  without being reflected by the jet boundaries.

TABLE 2

CALCULATION OF CHARACTERISTICS ON PHYSICAL PLANE

State	$\theta_*$ <sup>+</sup>	$\theta_*$ <sup>-</sup>	$\theta$	$v(M)$	M	$\mu$	$\theta + \mu$	$\theta - \mu$	P/P <sub>t</sub>
1	0	0	0	0	1	90°	90°	-90°	0.5283
2	0	10	10	10	1.435	44.15	54.15	-34.15	
3	0	20	20	20	1.775	34.3	54.3	-14.3	.2990
4	0	40	30	30	2.135	27.93	57.93	2.07	.1810
5	0	60	40	40	2.54	23.18	63.18	16.82	.1035
6	0	80	50	50	3.013	19.38	69.38	30.62	.55 -1
7	0	100	60	60	3.595	16.15	76.15	43.85	.2672 -1
8	0	120	67.7	67.7	4.145	13.96	88.7	53.74	.1146 -1
9	20	135.4	0	20	1.775	34.3	34.3	-34.3	.5437 -2
10	20	20	10	30	2.135	27.93	37.93	-17.93	.1810
11	20	40	20	40	2.54	23.18	43.18	- 3.18	.135
12	20	60	30	50	3.013	19.38	49.38	10.62	.55 -1
13	20	80	40	60	3.595	16.15	56.15	23.85	.2672 -1
14	20	100	50	70	4.339	13.315	63.32	36.69	.1146 -1
15	20	120	57.7	77.7	5.085	11.34	69.04	46.4	.425 -2
16	40	135.4	0	40	2.54	23.18	23.18	-23.18	
17	40	40	10	50	3.013	19.38	29.38	- 9.38	.55 -1
18	40	60	20	60	3.595	16.15	36.15	3.85	.2672 -1
19	40	80	30	70	4.339	13.315	43.32	16.69	.1146 -1
20	40	120	40	80	5.3479	10.777	50.78	29.23	.4215 -2
21	40	135.4°	47.7	87.7	6.433	8.943	56.64	38.76	
22	60	60	0	60	3.595	16.15	16.15	-16.15	.1146 -1
23	60	80	10	70	4.339	13.315	23.32	- 3.32	.4215 -2
24	60	100	20	80	5.3479	10.777	30.78	9.22	
25	60	120	30	90	6.8190	8.433	38.43	21.57	
26	60	135.4°	37.7	97.7	9.2105	6.2330	43.93	31.5	
27	80	80	0	80	5.3479	10.777	10.777	-10.77	
28	80	100	10	90	6.8190	8.433	18.433	1.57	
29	80	120	20	100	9.2105	6.233	26.23	13.8	
30	80	135.4	27.7	107.7	12.45	4.61	32.3	23.1	
31	100	100	0	100	9.2105	6.233	+ 6.233	- 6.233	
32	100	120	10	110	13.874	4.1331	14.13	5.9	
33	100	135.4	17.7	117.7	21.4	2.678	20.37	15.0	
34	120	120	0	120	27.335	2.097	2.097	- 2.097	
35	120	135.4	7.7	127.7	104	0.546	8.2	7.2	
36	-	-	11°	-	∞	0	-	-	0
51	20	115.4	47.7	67.7°	4.145	13.96		33.7	.5437 -2
52	40	115.4	37.7	77.7	5.085	11.34		26.36	
53	40	95.4	+27.7	67.7°	4.145	13.96		13.7	.5437 -2
54	60	75.4	+ 7.7	67.7°	4.145	13.96		- 6.26	.5437 -2
55	60	95.4	17.7	77.7	5.085	11.34		6.4	
56	60	115.4	27.7	87.7	6.433	8.943		18.8	

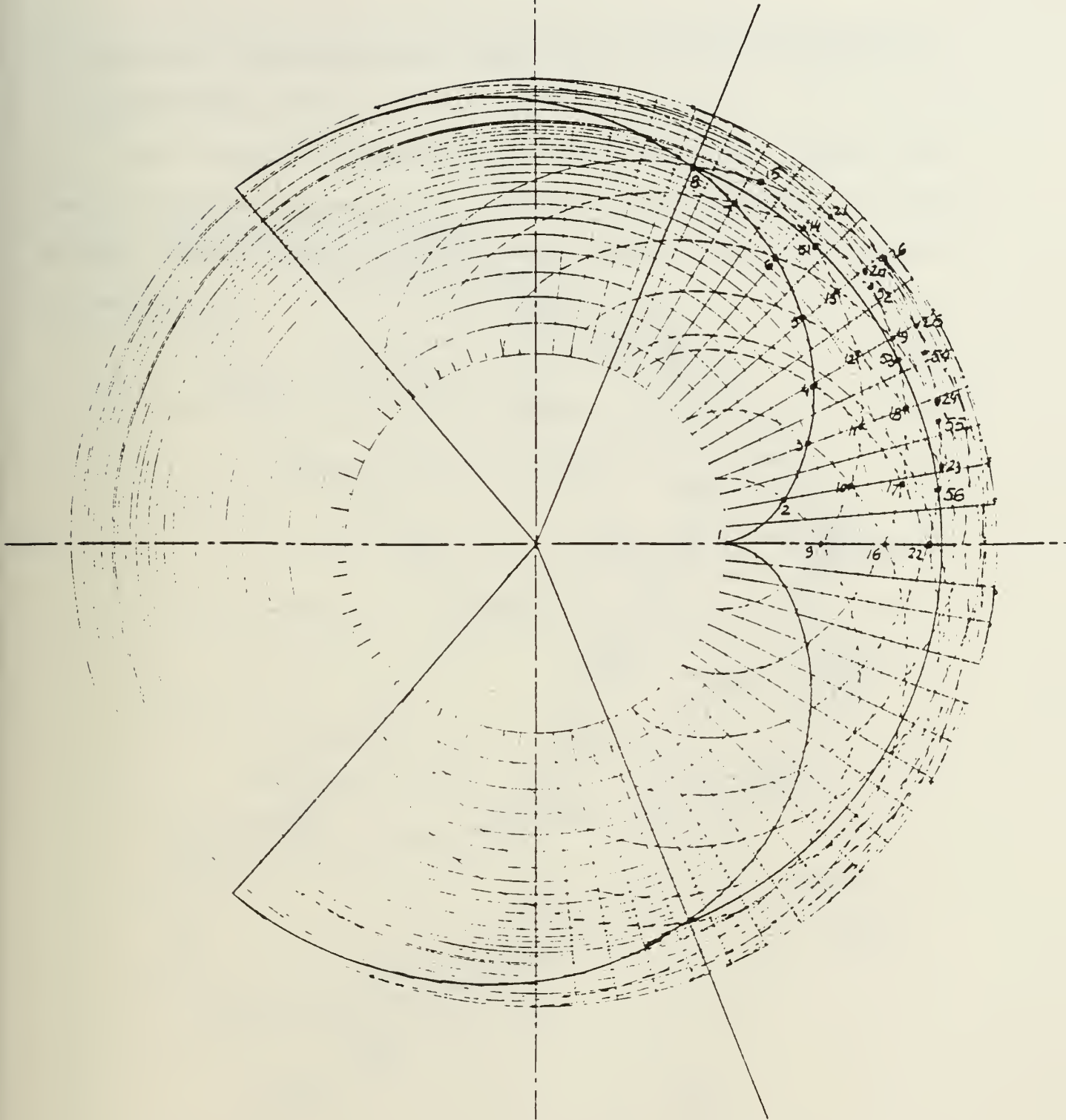


Figure 7. Hodograph plane showing the points on the mesh of characteristics.  
(Related to physical plan Figure 8).

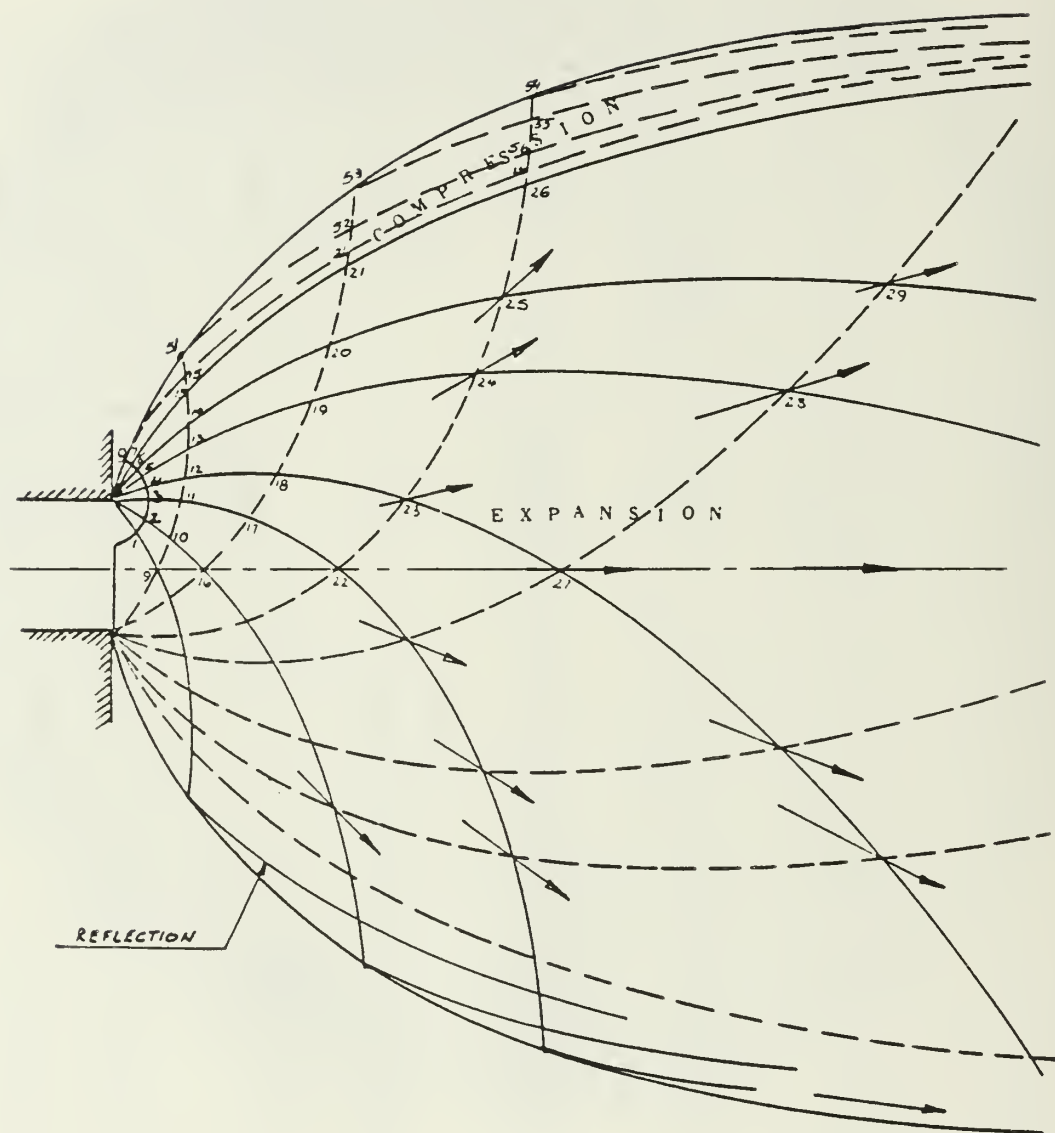


Figure 8. Schematic shape of a highly underexpanded jet.

(See Figure 7 - The Hodograph Plane)

The arrows indicate flow direction.

E. THE METHOD OF CHARACTERISTICS; COMPUTATION OF PLANAR AND AXISYMMETRIC TWO-DIMENSIONAL FLOWS

For a planar two dimensional flow, the Prandtl-Meyer function  $v$  and flow direction  $\theta$  at any point (3) in the field may be calculated using data of two other points (1) and (2) located on characteristic lines that intersect at (3) (see Fig. (9))

$$v_3 = \frac{1}{2} (v_1 + v_2) + \frac{1}{2} (\theta_1 - \theta_2) \quad (7)$$

$$\theta_3 = \frac{1}{2} (v_1 - v_2) + \frac{1}{2} (\theta_1 + \theta_2) \quad (8)$$

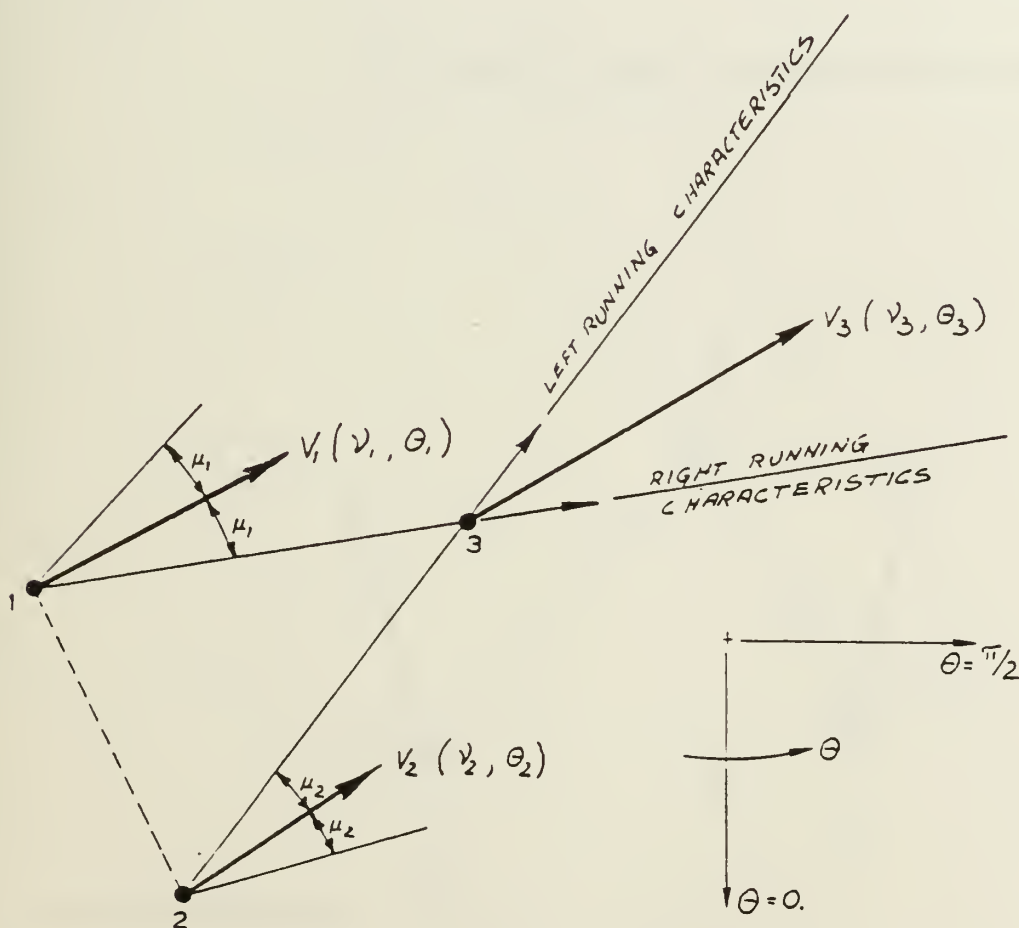


Figure 9. The calculation of  $v$  and  $\theta$  for point (3) is based on data for points 1 and 2.

For axisymmetric, two-dimensional flow, Liepmann and Roshko [2] developed expressions for finding the Prandtl-Meyer function ( $v$ ) and the flow direction ( $\theta$ ) which are given by:

$$v_3 = \frac{1}{2} (v_1 + v_2) + \frac{1}{2} (\theta_1 - \theta_2) + \frac{1}{2} \left[ \sin \mu_1 \frac{\sin \theta_1}{r_1} \Delta \xi_{13} + \sin \mu_2 \frac{\sin \theta_2}{r_2} \Delta \eta_{23} \right] \quad (9)$$

$$\theta_3 = \frac{1}{2} (v_1 - v_2) + \frac{1}{2} (\theta_1 + \theta_2) + \frac{1}{2} \left[ \sin \mu_1 \frac{\sin \theta_1}{r_1} \Delta \xi_{13} - \sin \mu_2 \frac{\sin \theta_2}{r_2} \Delta \eta_{23} \right] \quad (10)$$

The angles and subscripts are shown in Figure (10).

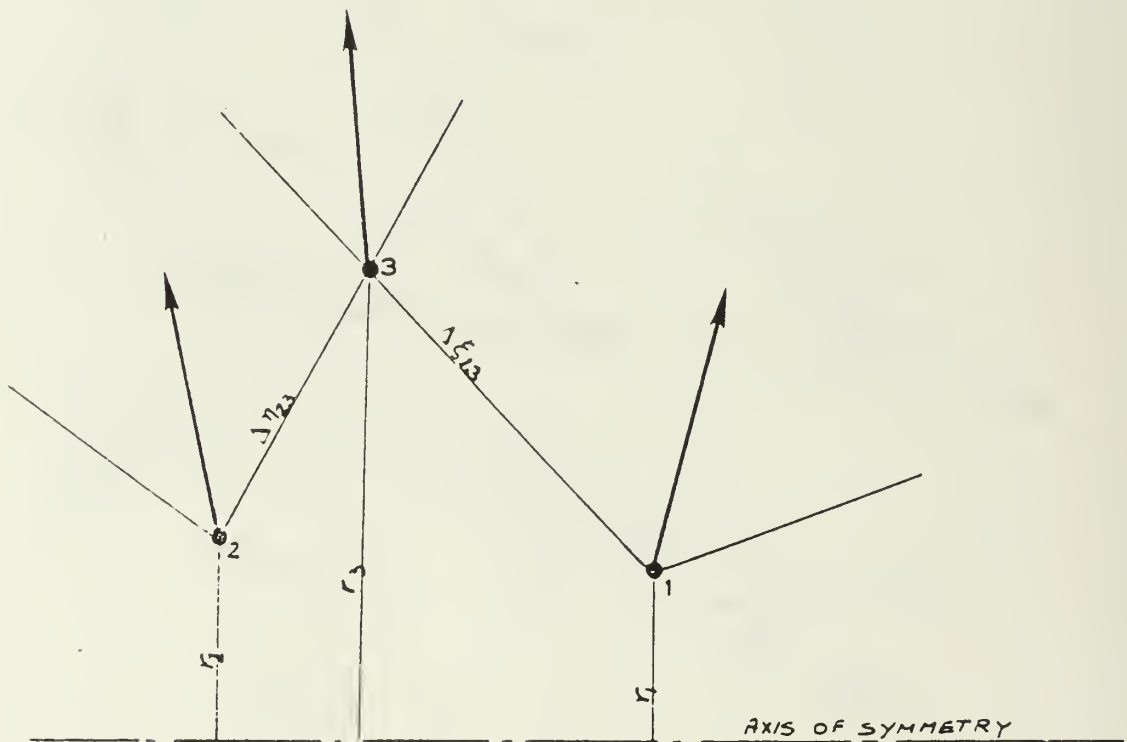


Figure 10. Calculation of  $\theta$  and  $v$  for axisymmetric flow.



It is obvious that for the axially symmetric flows the increase in the radius causes the increase in the flow cross section and influences the flow direction and the Prandtl-Meyer function. These facts have been taken into account when developing the "compatibility equations" (9,10).

Using equations (9,10) we have developed a computer program which enables the calculation of the jet flow for a two dimensional and for an axially symmetric geometry (ring jet).

The listing of the program, the program description and some results are given in Appendix A.

### III. THE BREAKDOWN OF THE CONTINUUM THEORY

#### A. GENERAL CRITERIA

As described in detail by Bird (Chapter 1 Reference [4]), the validity of the continuum approach has been identified with the validity of the Navier Stokes equations. This requires that the Knudsen number  $K_n = \lambda/L$  should be small compared with unity ( $\lambda$  is the mean free path and  $L$  is a scale length for the specific flow field). For  $K_n$  larger than a certain limit (between 0.01 to 0.1 depending on the required accuracy) a microscopic approach is necessary.

For small values of  $L$ , the microscopic approach may lead to statistical fluctuations of the results due to the small number of molecules participating in the flow processes. In figure (11) which was reproduced from Bird's book [4, figure 1.6], the regimes of rarefied flow and high fluctuations are depicted. The flow around the jet-air boundaries near a spacecraft is generally rarefied (high Knudsen number), but has insignificant fluctuations.

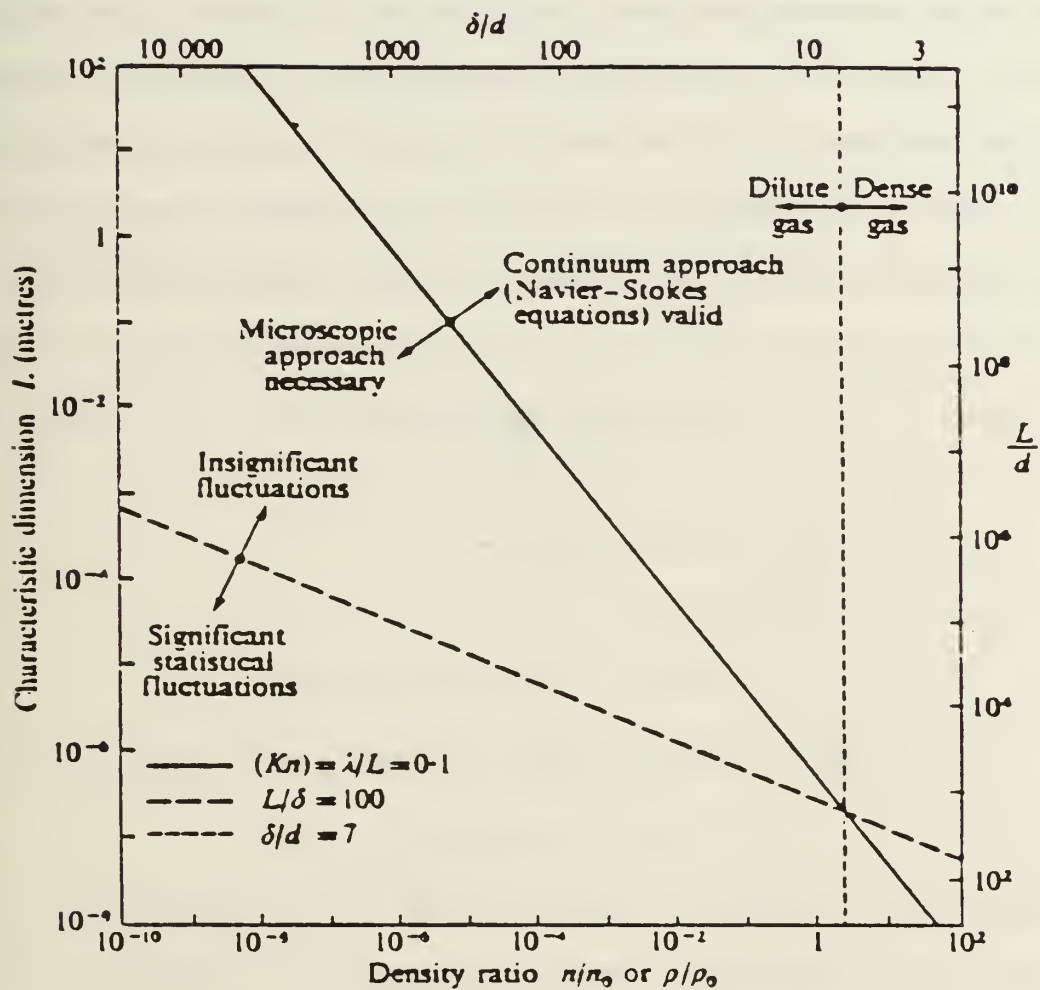


Figure 11. Limits for continuum approach and microscopic approach ( $d=3.7 \times 10^{-10}$  m).

## B. THE EMPIRICAL CRITERION

The Method of Characteristics (MOC) was used to compute the jet flow and the results obtained from the computer program "AXSYM" are valid as long as the continuum flow theory is valid.

The continuum flow requires that the mean free path should be negligibly small in comparison with the scale length of the macroscopic flow variations. The classical theory for Prandtl-Meyer expansion may therefore be expected to fail at progressively larger distances from the nozzle lips as the gas density decreases with the increasing flow angle and Mach number. The empirical criterion for the breakdown of continuum flow in steady expansion flow [4] is that

$$P \equiv \frac{q}{\rho v} \left| \frac{d\rho}{dS} \right| \approx 0.05 \quad (11)$$

where

$q$  = stream velocity

$\rho$  = density

$v$  = molecular collision frequency

$\left| \frac{d\rho}{dS} \right|$  = absolute change in density while moving a distance  
dS along a streamline

Introducing the breakdown parameter  $P$  into the program, gives the definition of the boundary where the flow should be calculated by means of the molecular flow theory, i.e., by solving the Boltzmann equation.

For an underexpanded jet with a high initial Mach number, the breakdown surface is nearly a streamline. Furthermore, the range of flow parameters for the present problem are such that the simple region extends to very large distances and near the nozzle lip the breakdown limit may be approximated by a straight line.

For the axisymmetric jet there is no simple region, however, for the region of interest it may be regarded as linear.

The method proposed in the present work for solving the flow behind the breakdown boundary is the Direct Simulation Monte Carlo (DSMC). For this purpose, a computer program "SIMUL" was developed. In the following chapter we describe the algorithms required for the specific problem, the geometry and the data organization. Detailed program description is given in Appendix (B).

#### IV. THE MOLECULAR FLOW IN AN AXISYMMETRIC RING JET

##### A. GENERAL CONSIDERATIONS

The part of the field in which the jet may be calculated by means of the continuum theory was described in Chapter II. There we calculated also the boundaries where continuum theory becomes invalid and molecular calculation should be employed. In fact, the molecular theory and the molecular Boltzmann equations are universal and hold for the entire flowfield. However, computational requirements make the Boltzmann equation impractical for the upstream flows. Therefore we limit our solution only to the part of the flow beyond the region where continuum breakdown occurs.

As a result obtained from MOC solution the "breakdown", i.e., the locus where the breakdown parameter  $p$  has values between 0.03 to 0.06, for the region close to the nozzle lips this boundary may be approximated by a straight line (for axisymmetric flow this line is the envelope of a cone, see Figure (12)).

For the specific jet and gas, the breakdown occurs in a region where the number density is in a range of  $10^{21}$  molecules/m<sup>3</sup>. For ambient gas at an altitude of 200 km the number density is  $10^{15}$  and decreases to a range of  $10^{11}$  at 1000 km. In order to be able to express this vast change in a simulation, we would need to have an extremely large number of cells which would be impossible to store in a computer. To overcome this problem we are required to make a less exact formulation which enables the production of results, having to pay the penalty of "smearing" the steep gradients and obtaining averages within layers of simulated cells. Unfortunately it is impossible to



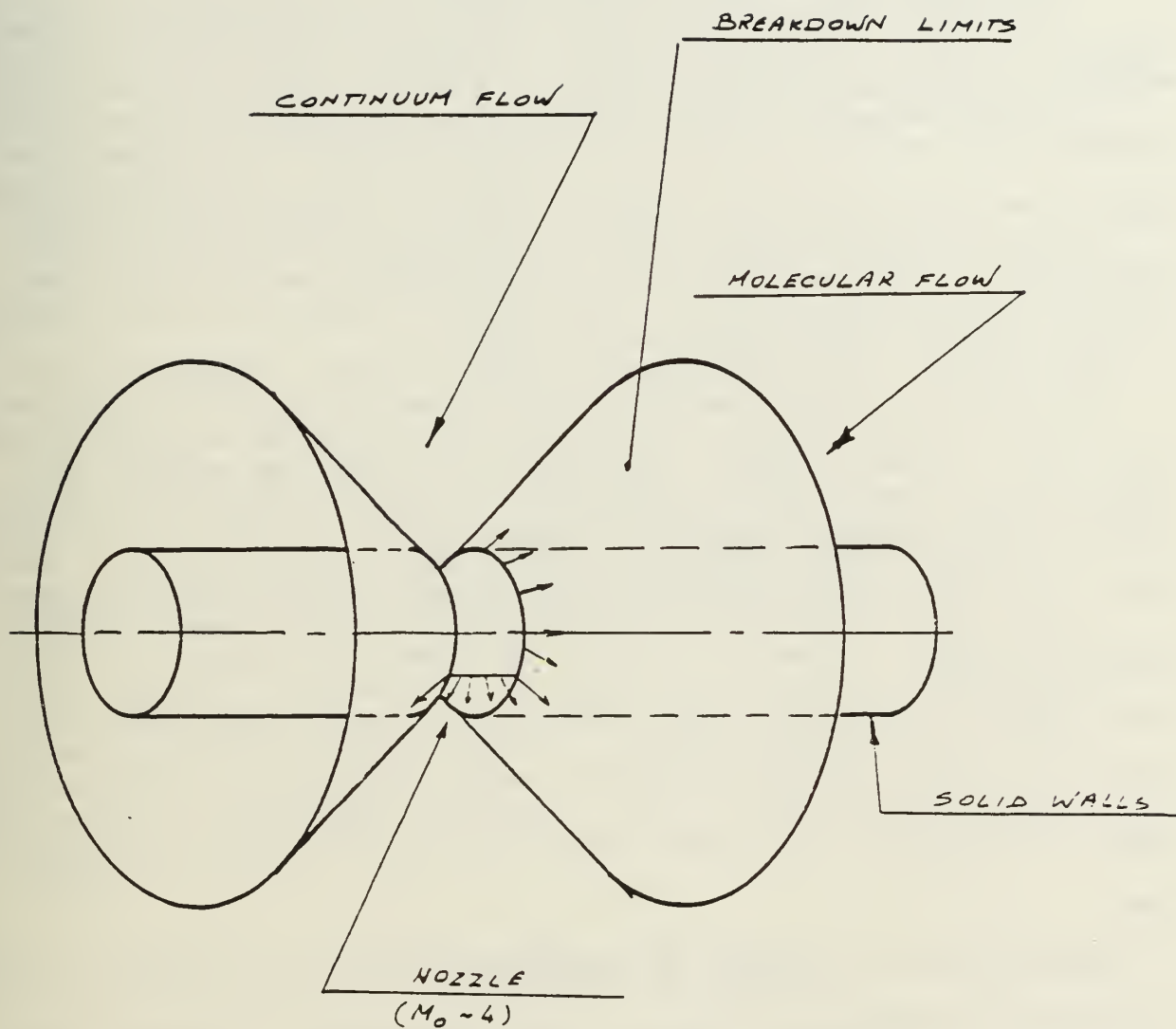


Figure 12. Regions in a ring jet.

predict how far the simulated results will be from the exact solutions. These comparisons have to be made after getting final results of this simulation.

### 1. The Direct Simulation Monte Carlo Method

The direct simulation Monte Carlo Method is a technique for a computer modeling of a real gas by some thousands of simulated molecules. The velocity components and the coordinates of the simulated molecules are stored in the computer and are modified with time as a result of collisions and boundary interactions. A detailed description of some problems and their solutions by means of direct simulation is given in [4].

To follow the molecular motion it is necessary to divide the simulated domain into a network of cells. The size of a cell must be such that the change in flow properties across each cell is small. The time is advanced in discrete steps DTM, such that DTM is small compared with the mean collision time per molecule. If there is a flow going through the domain, DTM should be small compared with the mean time required for the mean flow to cross the cells.\* Both cell size ((DR),(DDALFA) - radial size and angular size as they appear in the program) and DTM may vary in the simulation with position and time.

Applications such as free jet expansion in which large gradients of flow properties are expected, may require a very large number of cells for the simulation. In these cases the computer memory requirements to store cells' data and molecules' data may exceed the available computer storage. A

---

\*If DTM is chosen to be very small compared with the mean time between collisions then the simulation will require a very large number of runs such that the number of collisions will be sufficient. If DTM is large the molecules are washed out by the mean flux and there is no time for the collisions to influence the flow.

solution for this problem is to divide the simulation space into smaller regions and to run the simulation for each region separately. If there is an interaction between different regions, which apriori is undefined, the solution should be found iteratively. (That means that each run will provide data for consecutive runs and the procedure should be repeated until the results converge to a steady solution.

The computation of a representative set of collisions based on mean collision time per molecule is invalid for a computerized simulation because of the large computer time and computer memory requirements. Instead, the method proposed by Derzko which is described in details by Bird [4] may be employed. Following this method, an averaged mean time between collisions of species L with species M for a cell is calculated. The number of collisions of each type (L.M species) is such that the collision time counters are kept concurrent with the overall time parameter. The L.M collision time for a cell containing  $N_L$  and  $N_M$  molecules with collision cross section  $\sigma_{LM}$ , number densities  $n_L$  and  $n_M$  and relative velocity  $C_r$ , is given by

$$\Delta t_c = \frac{LP}{N_L} \frac{1}{\sigma_{LM} n_M C_r} + \frac{MP}{N_M} \frac{1}{\sigma_{LM} n_L C_r} \quad (12)$$

where LP and MP are the probabilities that the collision will be effective for the L and M molecules respectively.

## B. THE GEOMETRY OF THE SIMULATED DOMAIN, SECTORS, REGIONS AND CELLS

Figure (13) shows a cross section of the simulated domain for the axisymmetric (ring) flow. Points A and A' are the nozzle lips. Starting at "A" and assuming the "breakdown" boundary to be a straight line, we obtain the cross section of the molecular domain as a sector defined by LAM. The solid wall is defined by AL. The arc LM may be assumed to be far enough so that the pressure along it may be assumed to equal the ambient pressure. Molecules originated in the jet cross the breakdown boundary with a velocity, direction, temperature (and other thermodynamic properties) as found from the continuum solution.

The molecular domain LAM is divided into secondary sectors, and each of these are divided into several radial regions making the "simulation regions".

Because we have no apriori information on how the expansion occurs, the angle of each sector (which mainly is in the direction of the expansion gradients) is left to be a result of the internal calculation.

Each region is divided into NRD radial divisions and NAD angular divisions making a network of  $NAD \times NRD$  simulation cells. The angle DALFA of all regions in a sector is constant. Taking NAD constant for all regions in a sector, we get the angle of a cell DDALFA constant. Defining the radial size of a cell DR as constant we get a cell cross section area proportional to the radius R measured from the nozzle lip (point A).

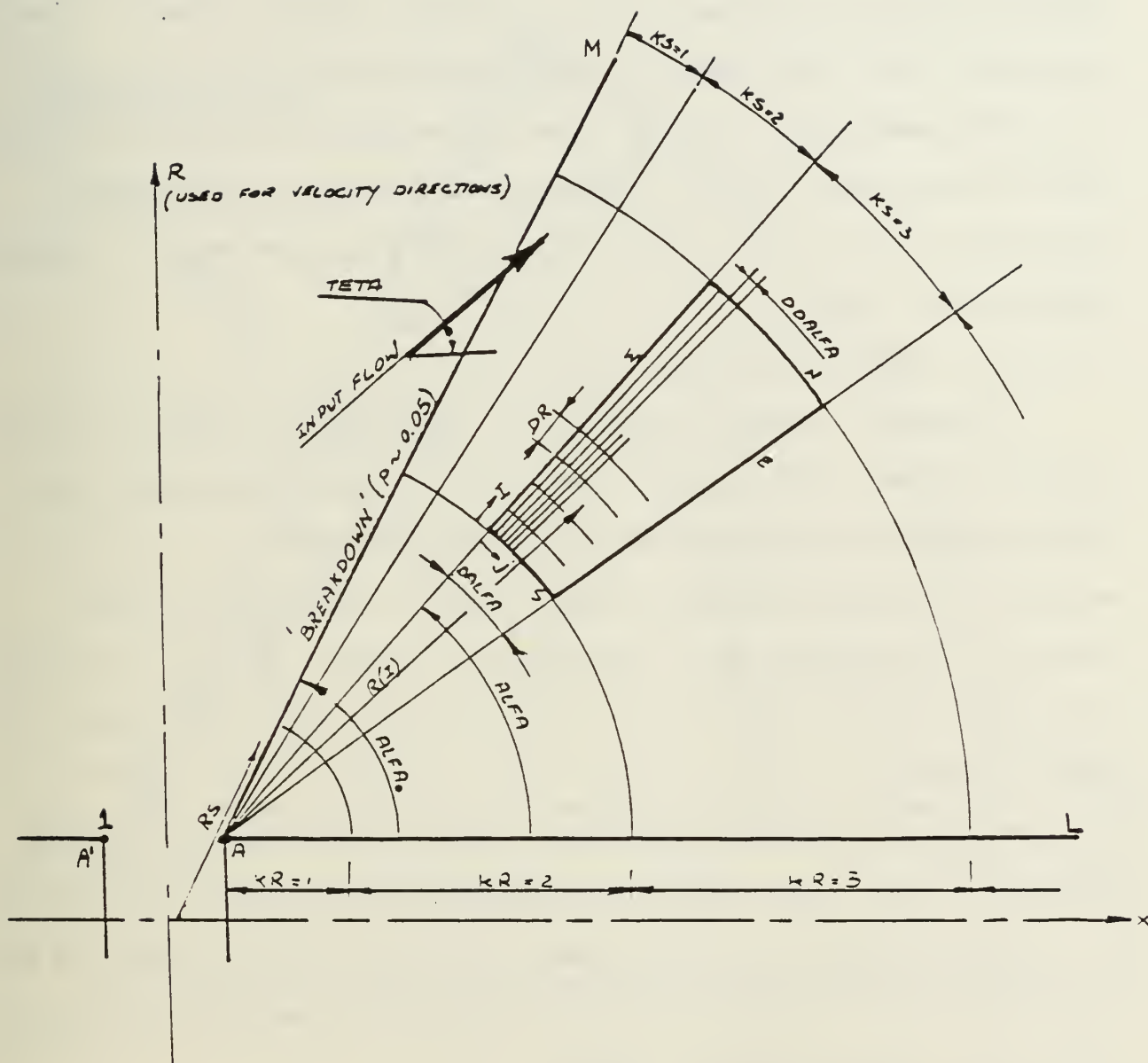


Figure 13. Cross section of the simulation domain, definition of sectors, regions, cells and coordinates.

The size of a cell: In order to get accurate simulated results it is recommended to define the size of a cell (DR and R\*DDALFA) small compared with the mean free path of the molecules  $\lambda$  (typical  $DR = \lambda/3$ ). However, as we do not expect to get large changes in flow parameters along the radius we may allow DR be much larger than  $\lambda/3$ . The angular size of the largest cell in a sector should comply with this requirement, but because of the computer limitation it is set to be equal to  $5*\lambda$ . This will be the basis for defining DALFA for each sector.

### C. INITIAL NUMBER OF MOLECULES IN CELLS

A "reasonable" number of molecules in a simulation is several thousands (a larger number, which is better, may be used for simple problems or when using a single user computer with large user memory space). The initial setting of molecules in cells is usually based on a guess of the number density in the specific cell. (The number of molecules in cells will change during the simulation according to the input/output calculated fluxes to the specific region).

The number density and the size of a cell are specified only in three dimensional flows. When applied to a two dimensional flow the simulation may be regarded as applying to an arbitrary thin slice of the real flow. In the axisymmetric flow we define the width of a cell by the angle DFI as shown in Figure (14), constant within a region.

The initial number density in cells of a given region is set constant. Defining the total number of simulated molecules in the region the number of molecules in each specific cell becomes a function of DFI. For example, assume that we limit the number of molecules in the smallest cell in the region to 15 then:



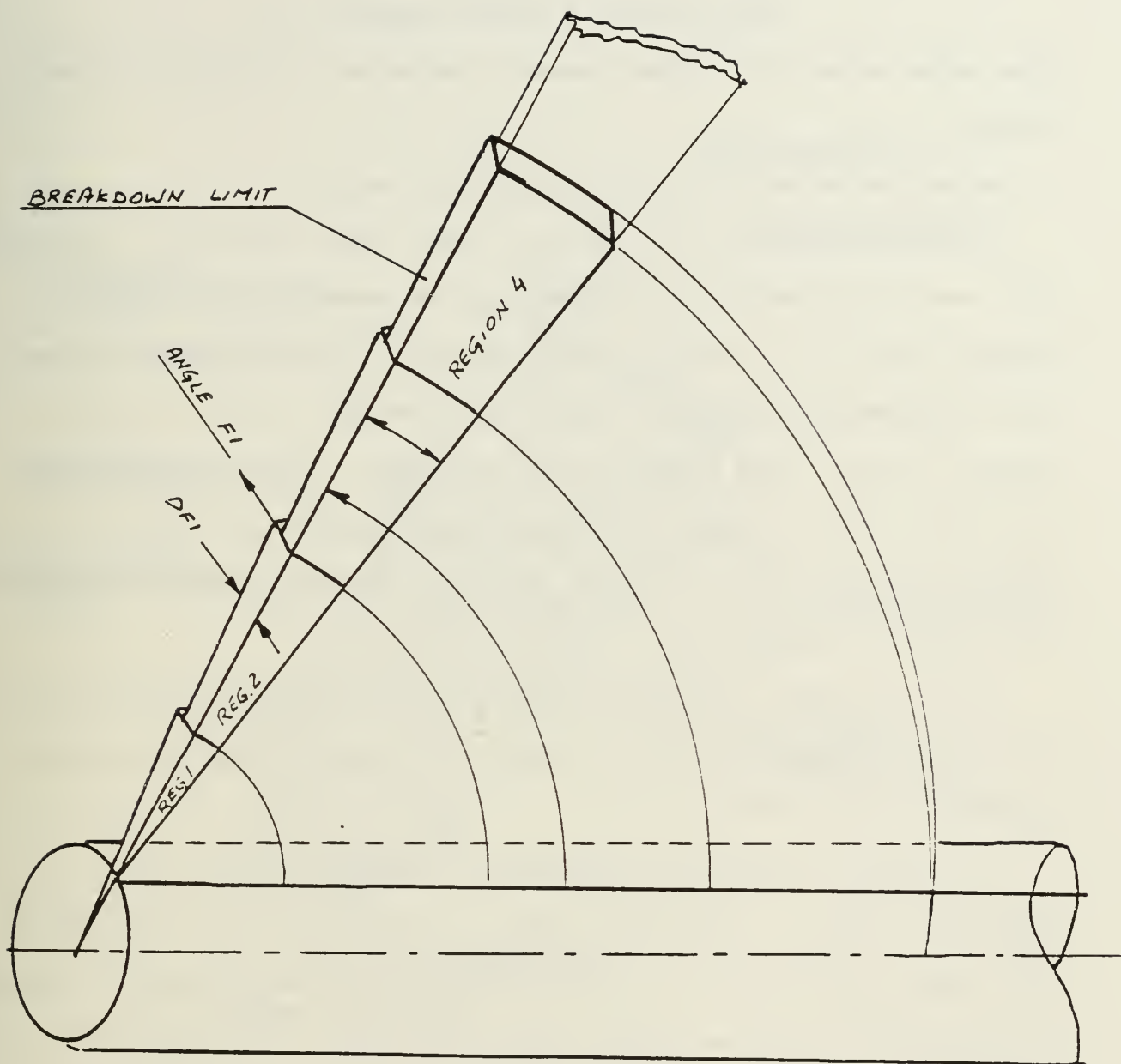


Figure 14. Variation of the angle DFI.

To maintain the number of simulated molecules within computational limits, the 'width' of each region defined by DFI is such that:

MIN = number of molecules in smallest cell in a region

MIN = VOLUME (smallest cell) \* number density

=  $f(R, ALFA, DALFA) * DFI * \text{number density for flux calculations}$

DFI is a weighting factor.

$$DFI*(contant) * (number\ density) = 15$$

Other cells contain the initial number of molecules proportional to their volumes.

#### D. DEFINITION OF INPUT AND OUTPUT FLOWS FOR A REGION

The cross sections of all regions (except those regions near the nozzle lips) are quadrilateral. Through the sides of the region molecules are allowed to enter or to leave according with the boundary conditions or as a result of molecular velocity. For the first sector, near the breakdown boundary the input flow (FWP1 and FWP2 see Figure (15)) is defined by the results from the continuum flow. FEN1, FNN1, FSN1, etc. are results of counting and averaging the outgoing molecules (the different vector names will be explained in Appendix B).

For the neighbor regions these output fluxes become inputs and have to be adjusted according to the differences in the angle DF1 of the different regions.

The simulation starts with regions in the sector near the breakdown boundary. At this time there is no data for input flows through faces E and N of the cell. An additional run of the whole program is required in order to take these calculated flows into account. If the accuracy of the results is important we may run this type of iteration several times until the results become stable. (Only after running the program for the whole domain once we shall be able to evaluate the importance of these iterations.)

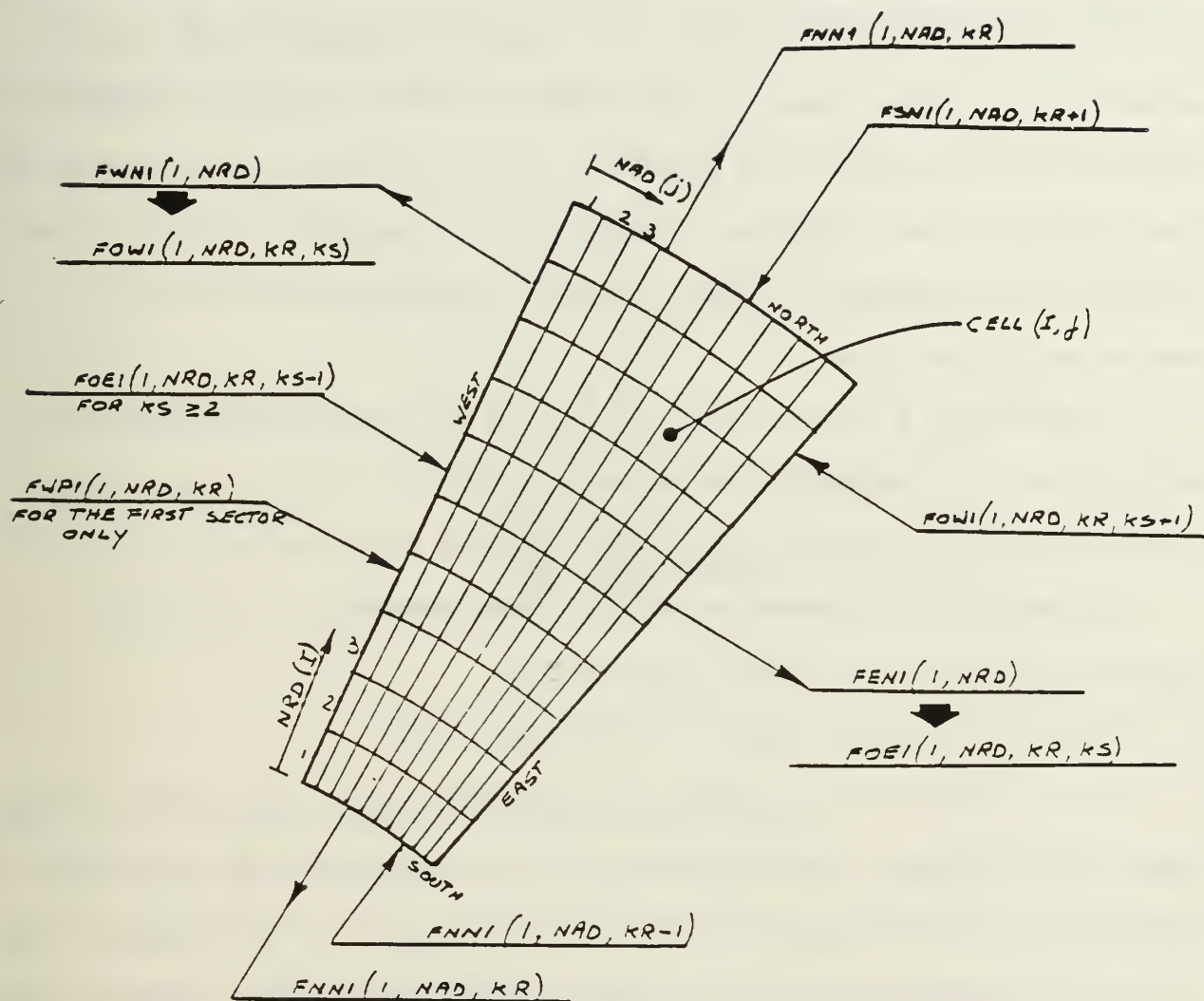


Figure 15. Definition of input and output flows of species 1 to a region (KR) in a sector (KS).

(For species 2 the flux names will change as follows:

instead  $FWN1( ) \rightarrow FWN2( )$

instead  $FOW1( ) \rightarrow FOW2( )$

etc.

## E. COLLISIONLESS FLOW

In several sectors near the breakdown boundary we may find a high number density and the mean free path small compared with the size of a cell. There the calculated collisions are expected to have an influence on the flow parameters. For wider expansion angles the collisions become rare mainly because of the decrease in the density. In the ambient gas the mean free path (for 200 km altitude) is 240 m. Comparing this number with the size of the simulated domain may lead to the conclusion that there the flow may be regarded as collisionless.

We may define a limiting line in the flow where the collisions become insignificant. Consequently, molecules crossing this limit will in fact continue moving in straight lines; a part of them reach the solid wall.

Introducing this idea of the collisionless flow we may reduce the computation time and the memory requirements.

## F. TWO DIMENSIONAL PLANAR FLOW VS. AXISYMMETRIC FLOW

The cell dimensions are completely specified only in three dimensional flow. When applied to two dimensional flow, the simulation may be regarded as applying to an arbitrarily thin slice of the real flow. The thickness of the slice may be chosen such that the number of simulated molecules complies with the cell volume and the physical number density. For the axisymmetric flow we have defined the angle DFI as the third coordinate so that the volume of the cell is completely specified.

Once the geometry is defined, the simulation may be accomplished and there is no difference if doing it for two dimensional or for axisymmetric flows.

## A.1 DIFFERENT REGIONS IN THE JET

For the two dimensional jet with initial Mach number greater than unity the different regions are shown in Figure (16).

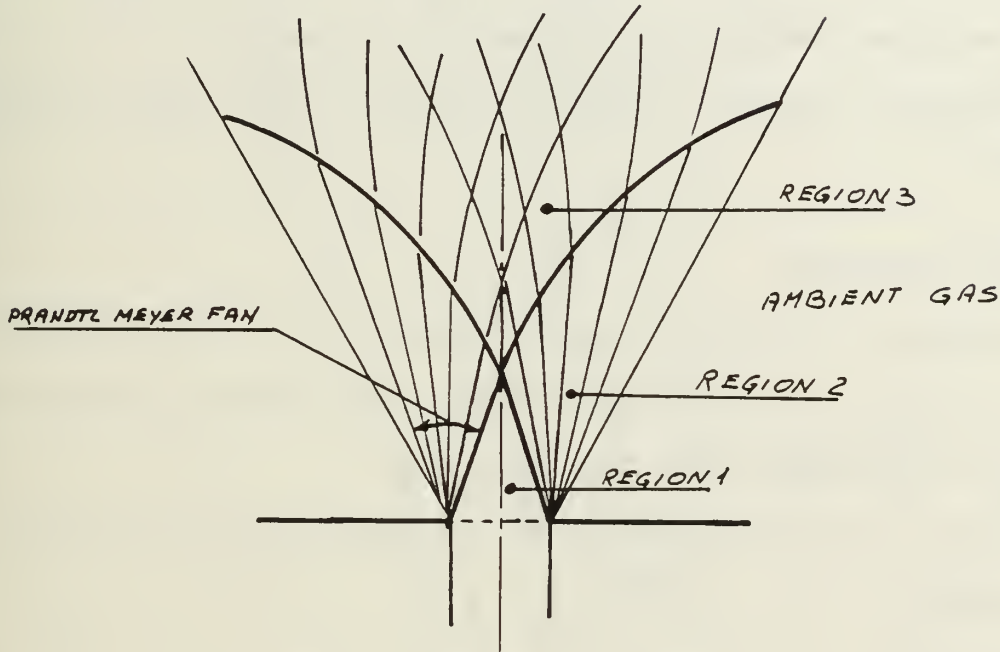


Figure 16. The three regions in an underexpanded jet.

For planar 2-D flow region 1 is a uniform flow core, region 2 is a simple region in which only one family of characteristics define the flow, and region 3 which contains the intersection of the two families of characteristics. Because our intention is to find solutions for highly underexpanded jets with very low ambient pressure, the calculation of further downstream flow is not necessary.

For the axisymmetric ring jet, we use the same definition for the different regions however, in this case none of the three regions has uniform flow and is not a simple region.



As shown in Equations (9,10) the PM function ( $\psi$ ) and the flow direction ( $\theta$ ) of a point at location  $I,j$  may be calculated from the  $\psi$  and  $\theta$  of two upstream points  $(I-1,j)$  and  $(I,j-1)$ . Later, from the PM function at the new point we may derive the local Mach number, the local pressure, temperature, velocity and other thermodynamic parameters as required.

Definition of the mesh of points for the different regions is shown in Figure (17).

## A.2 PROGRAM FLOWCHART

A simplified flowchart for the MOC program is shown in Figure (18). The program is designed to solve both axisymmetric as well as two dimensional flow

for  $kD = 2$  it solves two dimensional flow

for  $kD = 3$  it solves axisymmetric flow (This is also the default condition)

Initial data such as Mach number and pressure at the exit surface, ambient pressure and jet gas parameters are input data.

Output data contains the following for each mesh point:

Mach number, coordinates of mesh point  $(R,X)$ , flow direction (TETA), pressure, temperature, local velocity, Knudsen number based on the distance between two points along a streamline, mean free path and breakdown parameter as defined by Bird.\*

For each of the three regions, we start with precalculated boundary conditions enabling the calculation of the Mach angles, coordinates of mesh points and distances  $d\xi$  and  $d\eta$  as described in [2].

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\*For the exit plane instead the Bird's breakdown parameter, we calculate the ratio between time per three collisions and time of motion. Sometimes this ratio may be regarded as a measure of the breakdown of the continuum theory.



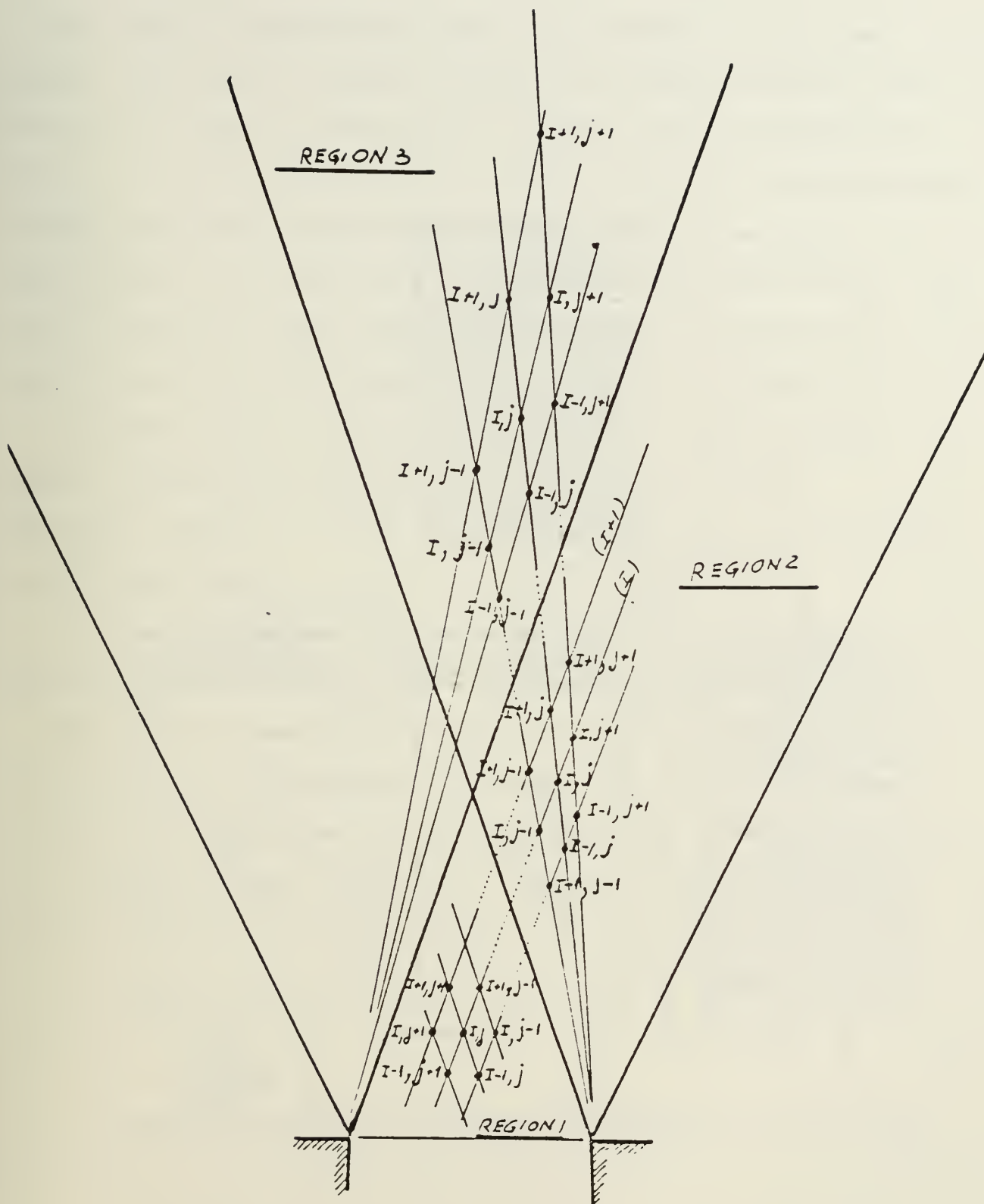


Figure 17. Indexing of mesh points for the different regions in the 'AXSYM' program.

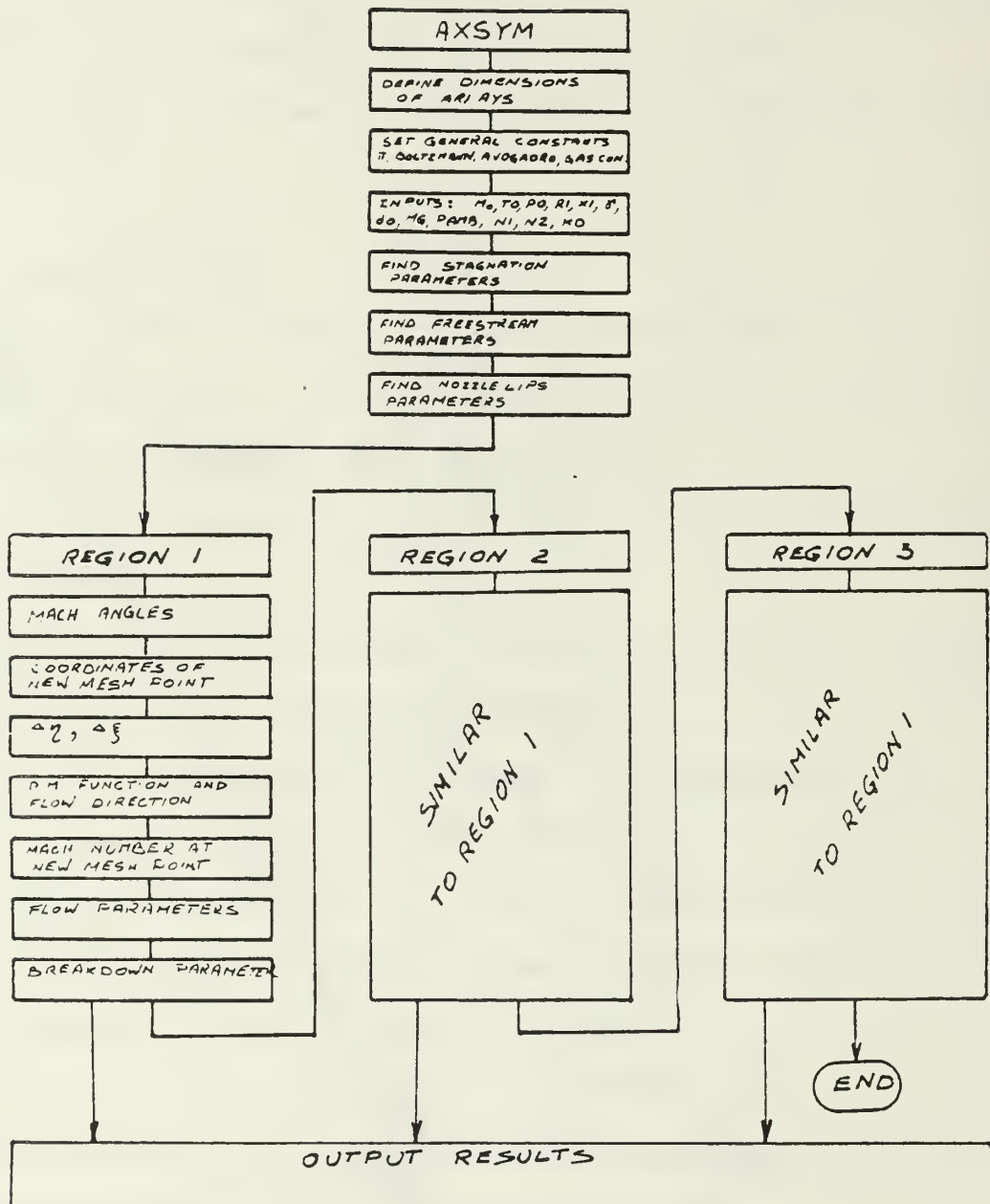


Figure 18. AXSYM Program Flowchart

The number of characteristics used in the program is arbitrary and depends on the required resolution (it may affect also the accuracy of the results and the amount of computation). We start with 20 characteristics along the exit cross section and with 50 characteristics in the Prandtl-Meyer fan thus a total of 70 characteristics of each family are calculated. In region 1 there are 20 left running and 20 right running characteristics. In region 2 there are 20 right running and 50 left running characteristics. In region 3 there are 50 left running and 50 right running characteristics.

In region 3 we limit the calculation where the two characteristics defining a new mesh point intersect at an angle smaller than the computational accuracy. In fact this occurs far downstream where continuum theory becomes invalid.

After defining the mesh geometry (successively) we calculate the Prandtl-Meyer function and flow direction, using equations (9,10).

The local Mach number is an implicit function of the Prandtl-Meyer angle. It is calculated by iterations with an initial guess of local Mach number set equal to a precalculated Mach number at an adjacent point, and the slope of the function as given by Equation (5). Figure (19) shows the iterative procedure for evaluating the Mach number at each mesh point.

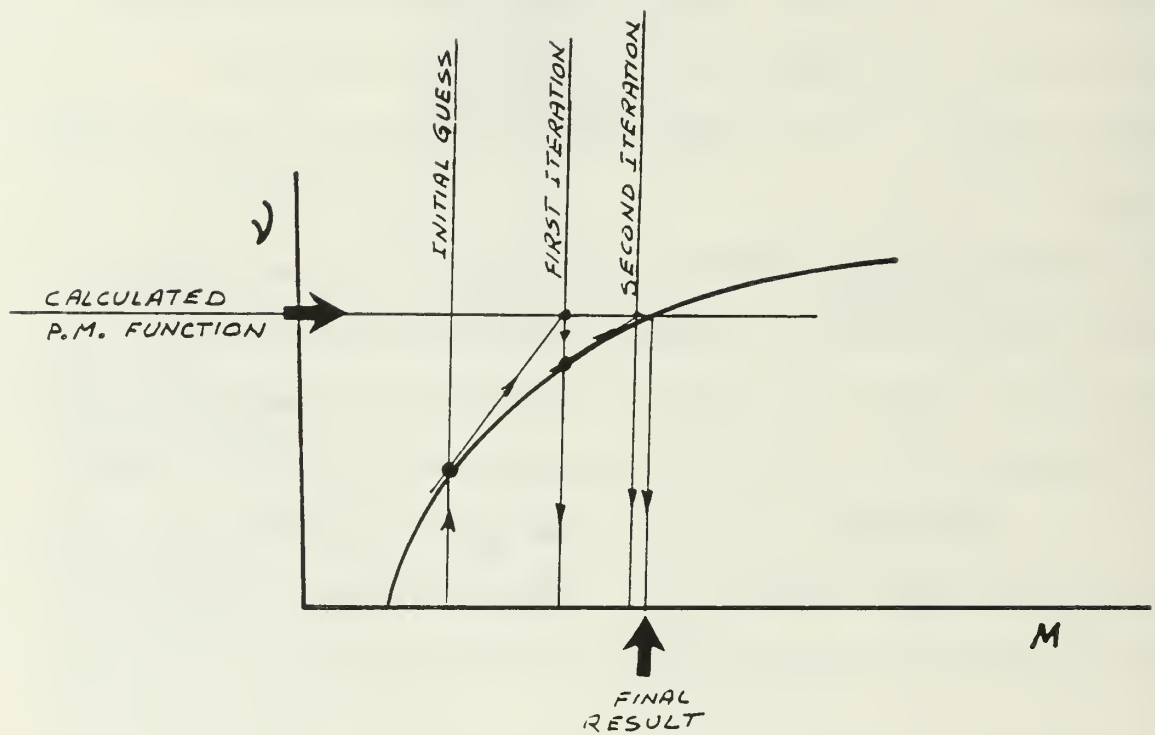


Figure 19. Iterative procedure for Mach number calculation.

Once the local Mach number has been found all other flow parameters may be defined using the Equation (1,2,3).

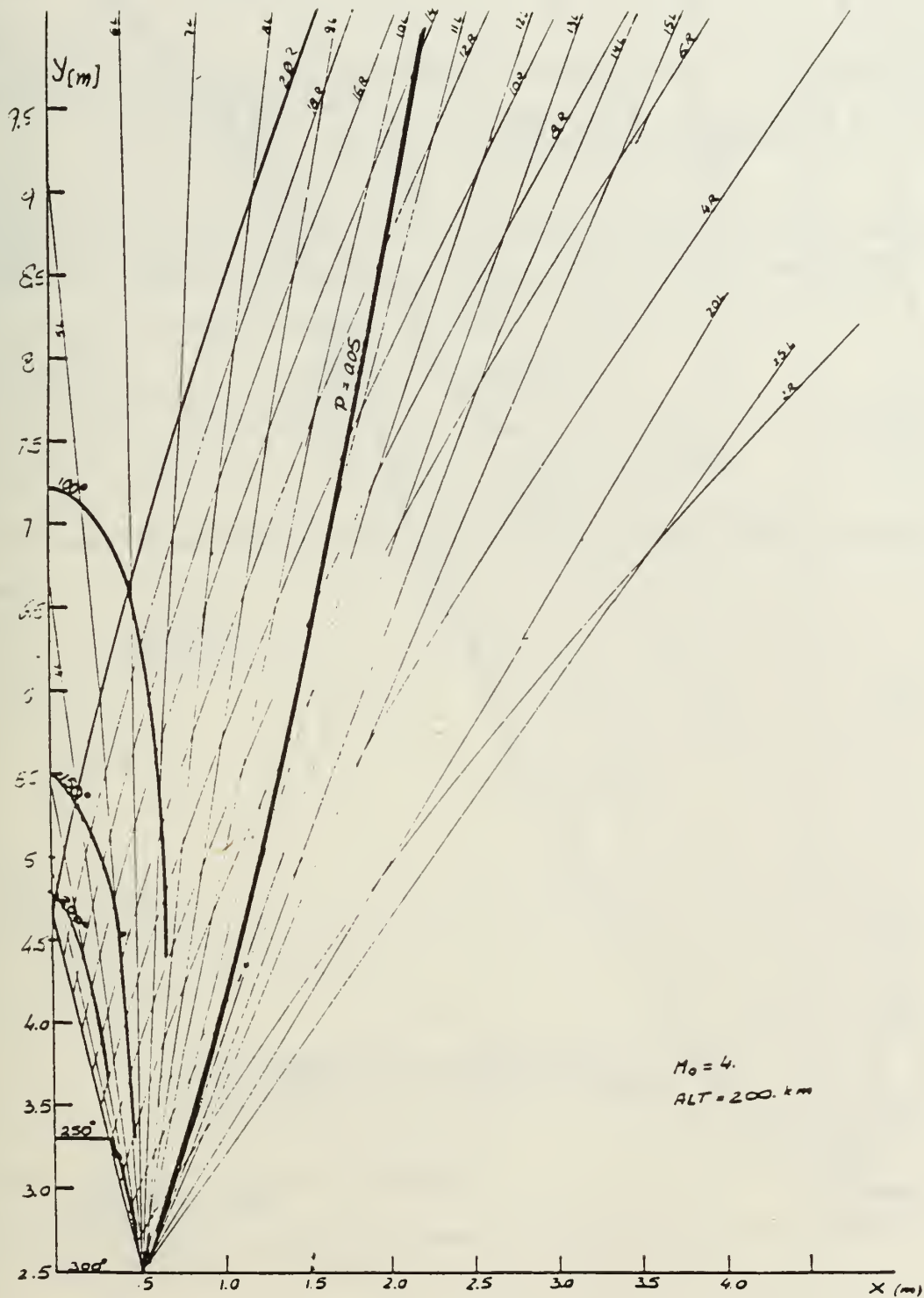


Figure 20. The mesh of characteristics in Region 2 and Region 3

Axisymmetric ring jet.  $M_0 = 4.$  Altitude=200km.

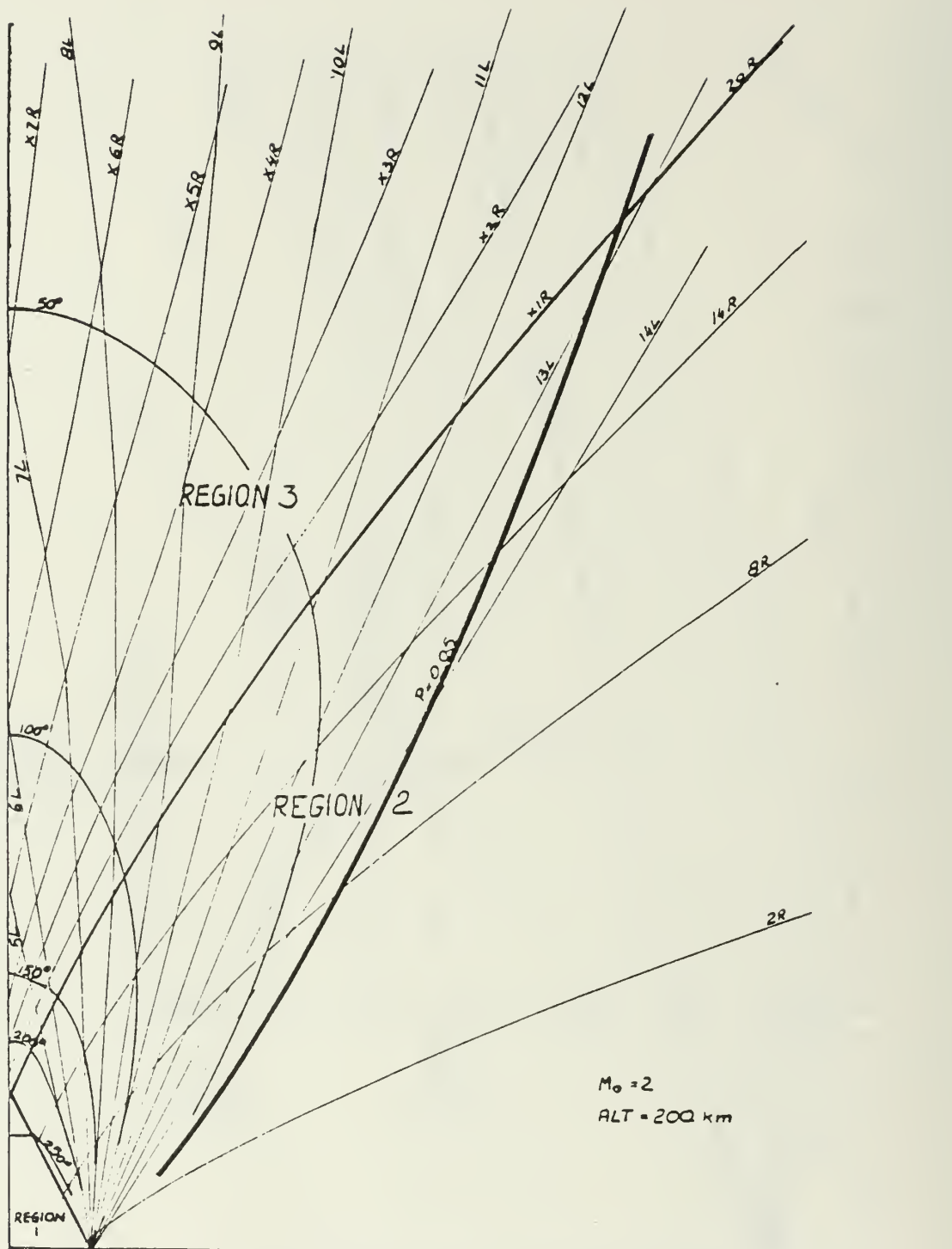


Figure 21. The mesh characteristics in Region 2 and Region 3  
Axisymmetric ring jet.  $M_0 = 2$ . Altitude = 200km.



### A.3 PROGRAM 'AXSYM' LISTING

```

$JOB
C PROGRAM AXSYM
C THIS PROGRAM CALCULATES THE ISENTROPIC EXPANSION OF A JET BY MEANS OF
C THE METHOD OF CHARACTERISTICS.
C FOR A TWO DIMENSIONAL JET 'KD' SHOULD BE SET EQUAL TO 2
C FOR AN AXISYMMETRIC RING JET 'KD' SHOULD BE SET EQUAL TO 3
C
C IMPLICIT REAL*8(A-H,O-Z,$)
C DIMENSION TETA(20,50),AM(20,50),R(20,50),X(20,50),PM(20,50)
C DIMENSION AMCOR(20,20),TETAC(20,20),XC(20,20),RC(20,20),PMC(20,20)
C DIMENSION DENSF(20,50)
C DIMENSION AMX(50,50),TETAX(50,50),XX(50,50),RX(50,50),PMX(50,50)
C
C TETA IS THE FLOW ANGLE (RADIAN) MEASURED FROM X AXIS.
C AM IS THE MACH NUMBER
C R IS THE RADIUS (NORMAL TO THE WALL)
C X IS THE AXIAL LOCATION (PARALLEL TO THE WALL)
C PM IS THE PRANDTL MEYER FUNCTION
C*****
C THE FOLLOWING IS DATA FOR THE SPECIFIC PROBLEM
C PAMB = 8.4736E-5
C KD = 3
C FOR TWO DIMENSIONAL FLOW KD=2 ,FOR AXISYMMETRICAL FLOW KD=3
C*****
C
C CONSTANTS
C PI = 3.141593
C BOLTZ = 1.38032E-23
C AVOG = 6.0225E+26
C RG = 8314.3
C
C EXIT SURFACE
C AM0 = 4.00
C T0 = 300.0
C P0 = 136.0
C
C
C R1 = 2.5
C X1 = 0.5
C
C GAS DATA
C GAMA = 1.535
C DIAM = 2.95E-10
C GM = 17.0
C RJ = RG/GM
C CXS = PI*DIAM*DIAM
C GMM = GM/AVOG
C
C MESH DEFINITION
C N1=CHARACTERISTICS FROM THE EXIT PLANE
C N2=CHARACTERISTICS FROM THE CORNERS
C
C N1 = 20
C N2 = 50
C
C CONSTANTS FOR THE ISENTROPIC EXPANSION
C A1 = (GAMA-1.0)/GAMA
C B1 = 1.0/A1
C
C A2 = DSQRT((GAMA+1.)/(GAMA-1.))
C B2 = 1.0/A2
C
C A3 = (GAMA-1.)/2.
C
C C = MACH*MACH*A3 + 1.
C D = MACH*MACH -1.
C*****
C*****
C DEFINE STAGNATION PARAMETERS
C
C C = (1.0+A3*AM0*AM0)
C PRESSURE

```

```

      PN = C**B1
      PSTG = P0*PN
C TEMPERATURE
      TSTG = T0*C
C DENSITY
      D0 = P0/(RJ*T0)
      DSTG = PSTG/(RJ*TSTG)
C
C*****
      WRITE(6,1)PSTG,TSTG,DSTG,DIAM,GM
      1 FORMAT('0','STAGNATION PRESSURE=',E12.5,' TEMPERATURE=',F10.5,
      1' DENSITY=',E12.5,' MOL.DIAM=',E12.5,' MOL.MASS=',F10.5)
      WRITE(6,2)P0,T0,D0,AM0
      2 FORMAT('0','EXIT PLANE PRESSURE=',E12.5,' TEMPERATURE=',F10.5,
      1' DENSITY=',E12.5,' MACH=',F10.5)
C*****
C*****
C DEFINE FREE STREAM PARAMETERS AT THE RIGHT CORNER
C
C*****
C MACH NUMBER
      FSM = DSQRT(((PSTG/PAMB)**A1-1.)/A3)
C TEMPERATURE
      FST = TSTG/(1.+A3*FSM*FSM)
C DENSITY
      FSD = PAMB/(RJ*FST)
C MACH ANGLES FOR HEAD AND TAIL OF FAN
      AMIT =DARSIN(1./AM0)
      AMIH =DARSIN(1./FSM)
C PRANDTL MEYER FUNCTION FOR HEAD AND TAIL OF FAN
      D1 =DSQRT(AM0*AM0-1.)
      D2 =DSQRT(FSM*FSM-1.)
      PMH = A2*DATAN(B2*D2)-DATAN(D2)
      PMT = A2*DATAN(B2*D1)-DATAN(D1)
C EXTERNAL TURNING ANGLE (FREE STREAM ANGLE)=EXTA
      EXTA = PMH - PMT
C PRANDTL MEYER FAN ANGLE PMFA
      PMFA = EXTA - AMIH + AMIT
C
C*****
C      WRITE(6,3)PAMB,FST,FSD,FSM
C      3 FORMAT('0','FREE STREAM PRESSURE=',E12.5,' TEMPERATURE=',F10.5,
C      1' DENSITY=',E12.5,' MACH=',F10.5///)
C*****
C*****
C DEFINE THE MESH OF CHARACTERISTICS-AND FLOW PARAMETERS
C
C*****
C THE CORNER POINT - LEFT RUNNING CHAR.
C PMFA IS DEVIDED INTO N2 (=50) NONEQUAL DIVISIONS
C
      RAT = (FSM/AM0-1)/0.02
      RAT2 = FLOAT(N2-1)
      E1 = DLOG(RAT)/DLOG(RAT2)
C
      WRITE(6,4)
      4 FORMAT('1','PRANDTL MEYER FAN LINE MACH
      1 P.M. ANGLE TETA'/)
      DO 20 N=2,N2
      EN2 = FLOAT(N-2)
      EN1 = FLOAT(N-1)
      AMB = AM0*(1.+0.02*(EN2)**E1)
      AMF = AM0*(1.+0.02*(EN1)**E1)
      DELM = AMF-AMB
      AM(1,N) = AMF
      D1 =DSQRT(AM(1,N)*AM(1,N)-1.)
      PM(1,N) = A2*DATAN(B2*D1)-DATAN(D1)
      AMI =DARSIN(1./AM(1,N))
      TETA(1,N) = PI/2.-(PM(1,N)-PMT)
C TETA IS THE FLOW ANGLE ON THE CHARACTERISTICS AT THE CORNER

```

```

      R(1,N) = R1
      X(1,N) = X1
      ALFAL = TETA(1,N)+AMI
C ALFAL IS THE ANGLE OF THE LEFT RUNNING CHARACTERISTICS
C
C PRESSURE, TEMPERATURE AND DENSITY VARIATION AT THE CORNER
      C = AM(1,N)*AM(1,N)*A3+1.
      PRES = PSTG/(C*B1)
      TEMP = TSTG/C
      DENSF(1,N) = PRES/(RJ*TEMP)
C
C*****
      WRITE(6,5) N,AM(1,N),PM(1,N),TETA(1,N)
      5 FORMAT(' ',1I5,3E20.5)
C*****
      20 CONTINUE
C
C*****
      WRITE(6,10)
      10 FORMAT('0',1I,1J,1MACH,1R,1X,1TETA,
1ITEMP,1PRESS,1VELOCITY,1KNUDSEN,1MFP,1ND,
1P')
C
C*****
C*****
C DEFINE THE PARAMETERS AT THE EXIT SURFACE (PLANE)
C
C*****
      DO 25 J = 1,N1
      AMCOR(1,J) = AM0
      D = DSQRT(AM0*AM0-1.)
      PMC(1,J) = A2*DATAN(B2*D)-DATAN(D)
      25 TETAC(1,J) = PI/2.
C
C THE EXIT PLANE IS DIVIDED INTO (N1-1) DIVISIONS (N1 POINTS)
      DO 30 J=1,20
      SOUND = DSQRT(GAMA*RJ*T0)
      VELO = AMCOR(1,J)*SOUND
      DNO = P0/(BOLTZ*T0)
      FPO = .707/(DNO*CXS)
      DX = X1*2./FLOAT(N1)
      AKNO = FPO/DX
C
      XC(1,J) = X1*(1.-FLOAT(J-1)/FLOAT(N1-1)*2.)
      CENTR = DABS(XC(1,J))
      IF(CENTR.LT.0.001) XC(1,J) = 0.
      RC(1,J) = R1
C
C.....
C      KZ=1*****
      IF (J.GT.1) GO TO 29
C AT THE EXIT PLANE THE BREAKDOWN PARAMETER IS EVALUATED BY MEANS OF
C THE RATIO OF THE COLLISION TIME AND THE FLOW TIME. LENGTH SCALE IS
C THE MESH DIMENSION.
      SCALE = DX*DTAN(DARSIN(1./AM0))/2.
      TIME1 = SCALE/VELO
      TIME2 = 3./(4.*CXS*DNO*DSQRT(BOLTZ*T0/(PI*GMM)))
      P = TIME2/TIME1
      DENSF(1,1) = D0
      29 CONTINUE
C
      WRITE(6,11)1,J,AMCOR(1,J),R1,XC(1,J),TETAC(1,J),T0,P0,VELO,AKNO,FP
      10,DNO,P
      11 FORMAT(' ',2I4,5F10.3,3E12.3,F9.4,2E12.3)
      30 CONTINUE
C
C*****
C*****

```



C		AXS0217
C	CALCULATE THE FLOW PARAMETERS IN THE CORE BOUNDED BY THE TWO MACH	AXS0218
C	WAVES STARTING AT THE NOZZLE LIPS (CORNER POINTS)	AXS0219
C	AT THE EXIT THE FLOW IS ASSUMED TO BE UNIFORM	AXS0220
C		AXS0221
C	*****	AXS0222
	WRITE (6,198)	AXS0223
	198 FORMAT('1',' CORE '/')	AXS0224
	APM=0.	AXS0225
	BPM=0.	AXS0226
	CPM=0.	AXS0227
C		AXS0228
	DO 199 I = 2,N1	AXS0229
	WRITE (6,10)	AXS0230
	DO 199 J = 1,N1	AXS0231
	IF ((I+J-1).GT.N1) GO TO 199	AXS0232
	AMIL = DARSIN(1./AMCOR(I-1,J))	AXS0233
	ALFAL = PI-(TETAC(I-1,J)+AMIL)	AXS0234
C		AXS0235
	AMIR = DARSIN(1./AMCOR(I-1,J+1))	AXS0236
	ALFAR = TETAC(I-1,J+1)-AMIR	AXS0237
C		AXS0238
	XC(I,J) = (RC(I-1,J)-RC(I-1,J+1)+XC(I-1,J)*DTAN(ALFAL)+XC(I-1,J+1)	AXS0239
	1*DTAN(ALFAR))/(DTAN(ALFAL)+DTAN(ALFAR))	AXS0240
	CENTR = DABS(XC(I,J))	AXS0241
	IF (CENTR.LT.0.001) XC(I,J) = 0.	AXS0242
	RC(I,J) = RC(I-1,J+1)+(XC(I,J)-XC(I-1,J+1))*DTAN(ALFAR)	AXS0243
C		AXS0244
	DKSI = DSQRT((XC(I,J)-XC(I-1,J+1))*2+(RC(I,J)-RC(I-1,J+1))*2)	AXS0245
	DETA = DSQRT((XC(I,J)-XC(I-1,J))*2+(RC(I,J)-RC(I-1,J))*2)	AXS0246
C		AXS0247
C		AXS0248
C	CALCULATE NOW THE PRANDTL MEYER FUNCTION AND FLOW ANGLE IN CORE.	AXS0249
	APM = PMC(I-1,J)+PMC(I-1,J+1)+TETAC(I-1,J+1)-TETAC(I-1,J)	AXS0250
	IF (KD.EQ.2) GO TO 151	AXS0251
	BPM = DSIN(AMIR)*DSIN(TETAC(I-1,J+1))/RC(I-1,J+1)*DKSI	AXS0252
	CPM = DSIN(AMIL)*DSIN(TETAC(I-1,J))/RC(I-1,J)*DETA	AXS0253
	151 PMC(I,J) = (APM+BPM+CPM)/2.0	AXS0254
C		AXS0255
	APM = PMC(I-1,J+1)-PMC(I-1,J)+TETAC(I-1,J+1)+TETAC(I-1,J)	AXS0256
	TETAC(I,J) = (APM+BPM-CPM)/2.0	AXS0257
C		AXS0258
C	DEFINE MACH NUMBER (BY ITERATIONS)	AXS0259
C	INITIAL GUESS AMCOR(I,J)=AMCOR(I-1,J+1)	AXS0260
C		AXS0261
	AMG = AMCOR(I-1,J+1)	AXS0262
	KZ = 0	AXS0263
	154 IF (KZ.GE.100) GO TO 160	AXS0264
	KZ = KZ+1	AXS0265
	C = AMG*AMG*A3+1.	AXS0266
	D = DSQRT(AMG*AMG-1.)	AXS0267
	PMCAL = A2*DATAN(B2*D)-DATAN(D)	AXS0268
	DELNI = PMCAL - PMC(I,J)	AXS0269
	DEL = DABS(DELNI)	AXS0270
	IF (DEL.LT..000002) GO TO 160	AXS0271
	IF (DELNI.LT.0.) GO TO 156	AXS0272
	AMG = AMG*.999	AXS0273
	GO TO 154	AXS0274
	156 AMG = AMG*(1.-DELNI*C/D)	AXS0275
	GO TO 154	AXS0276
	160 AMCOR(I,J) = AMG	AXS0277
C	.....	AXS0278
C		AXS0279
C	CALCULATE FLOW PARAMETERS	AXS0280
	IF (J.GT.1) GO TO 197	AXS0281
	C = AMCOR(I,J)*AMCOR(I,J)*A3+1.	AXS0282
	PRES = PSTG/(C*B1)	AXS0283
	TEMP = TSTG/C	AXS0284
	DN = PRES/(BOLTZ*TEMP)	AXS0285
	FP = .707/(DN*CXS)	AXS0286
	SCALE = DKSI * DSIN(ALFAL)	AXS0287
	AKN = FP/SCALE	AXS0288

```

      SOUND = DSQRT(GAMA*RJ*TEMP)
      VEL = SOUND*AMCOR(I,J)
C
C BREAKDOWN PARAMETER AS DEFINED BY 'BIRD'.
      DENSF(I,J) = PRES/(RJ*TEMP)
      DDENS = DENSF(I-1,J) - DENSF(I,J)
      COLF = 4.*CXSDN*DSQRT(BOLTZ*TEMP/(PI*GMM))
      P = VEL*DDENS/(SCALE*DENSF(I,J)*COLF)
197 CONTINUE
C
C      TIME1 = SCALE/VEL
C      TIME2 = 3./(4.*CXSDN*DSQRT(BOLTZ*TEMP/(PI*GMM)))
C      P = TIME2/TIME1
C*****
C*****
C*****
      WRITE(6,11)I,J,AMCOR(I,J),RC(I,J),XC(I,J),TETAC(I,J),TEMP,PRES,VEL
1,AKN,FP,DN,P
199 CONTINUE
C
C*****
C*****
C MATCH CORE AND FAN POINTS
      DO 200 I=1,N1
        X(I,1) = XC(I,1)
        R(I,1) = RC(I,1)
        AM(I,1) = AMCOR(I,1)
        PM(I,1) = PMC(I,1)
      200 TETA(I,1) = TETAC(I,1)
C
C*****
C*****
C CALCULATE FLOW PARAMETERS IN REGION 2 (SIMPLE PRANDTL MEYER FAN).
C
C*****
      WRITE (6,298)
298 FORMAT ('1', '      REGION      2'//)
C
      KFIN = 51
C
      DO 299 I = 2,N1
        WRITE(6,10)
        DO 299 J = 2,N2
          IF (J.GT.KFIN) GO TO 299
          KZ = 0
          AMIL = DARSIN(1./AM(I-1,J))
          AMIR = DARSIN(1./AM(I,J-1))
          ALFAL = PI-(TETA(I-1,J)+AMIL)
          ALFAR = TETA(I,J-1)-AMIR
C
C CHECK ANGLES AND CALCULATE COORDINATES
          ANGLE1 = PI/2.-.000001
          ANGLE2 = PI/2+.000001
          IF (ALFAL.LE.ANGLE1.OR.ALFAL.GE.ANGLE2) GO TO 201
          X(I,J) = X(I-1,J)
          GO TO 207
201 IF (ALFAR.LE.ANGLE1.OR.ALFAR.GE.ANGLE2) GO TO 205
          X(I,J) = X(I,J-1)
          GO TO 207
C
205 X(I,J) = (R(I-1,J)-R(I,J-1)+X(I,J-1)*DTAN(ALFAR)+X(I-1,J)*DTAN(ALF
          IAL))/(DTAN(ALFAL)+DTAN(ALFAR))
C
207 IF (ALFAL.LE.0.000001.OR.ALFAL.GE.0.000001) GO TO 209
          R(I,J) = R(I-1,J)
          GO TO 213
209 IF (ALFAR.LE.0.000001.OR.ALFAR.GE.0.000001) GO TO 211
          R(I,J) = R(I,J-1)
          GO TO 213
C
211 R(I,J) = (X(I,J)-X(I,J-1))*DTAN(ALFAR)+R(I,J-1)

```

```

213 DKSI = DSQRT((R(I,J)-R(I,J-1))*2+(X(I,J)-X(I,J-1))*2)
DETA = DSQRT((R(I,J)-R(I-1,J))*2+(X(I,J)-X(I-1,J))*2)
C
IF (R(I,J).GT.0..AND.X(I,J).GT.0.) GO TO 219
KFIN = J-1
WRITE(6,12)
12 FORMAT(' ',' FURTHER POINTS ON THE CHARACTERISTICS ARE DIVERGENT')
219 CONTINUE
C LOCATION OF THE NEW MESH POINT HAS BEEN FOUND
C
C
C CALCULATE NOW PRANDTL MEYER FUNCTION AND FLOW DIRECTION FOR NEW POINT
APM = PM(I,J-1)+PM(I-1,J)-TETA(I-1,J)+TETA(I,J-1)
IF (KD.EQ.2) GO TO 251
BPM = DSIN(AMIR)*DSIN(TETA(I,J-1))/R(I,J-1)*DKSI
CPM = DSIN(AMIL)*DSIN(TETA(I-1,J))/R(I-1,J)*DETA
251 PM(I,J) = .5*(APM+BPM+CPM)
C
APM = PM(I,J-1)-PM(I-1,J)+TETA(I,J-1)+TETA(I-1,J)
TETA(I,J) = .5*(APM+BPM-CPM)
C
C CALCULATE NOW MACH NUMBER FOR EACH POINT
C INITIAL GUESS AM(I,J) = AM(I-1,J)
AMG = AM(I-1,J)
KZ = 0
254 IF (AMG.GT.200.) GO TO 257
D = DSQRT(AMG*AMG-1.)
GO TO 258
257 D = AMG
258 IF (KZ.GE.100) GO TO 260
KZ = KZ + 1
PMCAL = A2*DATAN(B2*D)-DATAN(D)
DELNI = PMCAL - PM(I,J)
DEL = DABS(DELNI)
IF (DEL.LT..000002) GO TO 260
IF (DELNI.LT.0.)GO TO 256
AMG = AMG*.999
GO TO 254
C//////////
256 IF (AMG.LT.2000.)GO TO 2560
KFIN = J-1
GO TO 299
C//////////
2560 D1 = (A3*AMG*AMG+1.)
AMG = AMG*(1.-DELNI*D1/D)
GO TO 254
C
260 AM(I,J) = AMG
C CALCULATE NOW LOCAL TEMPERATURE,PRESSURE,VELOCITY,KNUDSEN NO.
C = AM(I,J)*AM(I,J)*A3+1
PRES = PSTG/(C*B1)
TEMP = TSTG/C
DN = PRES/(BOLTZ*TEMP)
DENSF(I,J) = PRES/(RJ*TEMP)
DDENS = DENSF(I-1,J-1)-DENSF(I,J)
SCALE = DSQRT((X(I,J)-X(I-1,J-1))*2+(R(I,J)-R(I-1,J-1))*2)
C*****
C
271 FP = .707/(DN*CXS)
C
AKN = FP/SCALE
SOUND =DSQRT(GAMA*RJ*TEMP)
VEL = AM(I,J)*SOUND
C
C
COLF = 4.*CXS*DN*DSQRT(BOLTZ*TEMP/(PI*GMM))
P = VEL*DDENS/(SCALE*DENSF(I,J)*COLF)
C
C PRINT RESULTS FOR MESH POINTS
KL = (-1)*I
IF (KL.LT.0) GO TO 299
WRITE(6,11)I,J,AM(I,J),R(I,J),X(I,J),TETA(I,J),TEMP,PRES,VEL,AKN,

```



```

      1FP,DN,P
      299 CONTINUE
C
C*****
C*****
C MATCH 'REGION 2' AND 'REGION 3' POINTS
C
C*****
      L = KFIN- 1
      DO 300 J = 1,L
        XX(1,J) = X(20,J)
        RX(1,J) = R(20,J)
        AMX(1,J) = AM(20,J)
        PMX(1,J) = PM(20,J)
      300 TETAX(1,J) = TETA(20,J)
C*****
C*****
C
C CALCULATE FLOW PARAMETERS FOR REGION 3
C
C*****
      WRITE (6,397)
397  FORMAT ('1',' REGION 3' /)
      DO 399 I = 2,L
        DO 399 J = 1,L
          IF (J.GT.KFIN) GO TO 399
          IF (J.GT.I)GO TO 320
          WRITE (6,10)
320  KZ = 0
C
      AMIL = DARSIN(1./AMX(I-1,J))
      ALFAL = PI-(TETAX(I-1,J)+AMIL)
      IF (J.GT.I) GO TO 301
      ALFAR = ALFAL
      TETAX(I,J) = PI*.5
      TETAX(I,J-1) = PI-TETAX(I-1,J)
      XX(I,J) = 0.
      RX(I,J) = RX(I-1,J)+XX(I-1,J)*DTAN(ALFAL)
      RX(I,J-1) = RX(I-1,J)
      XX(I,J-1) = -XX(I-1,J)
      DKSI = (RX(I,J)-RX(I-1,J))/DSIN(ALFAL)
      DETA = DKSI
      PMX(I,J-1) = PMX(I-1,J)
      GO TO 316
301  AMIR = DARSIN(1./AMX(I,J-1))
      ALFAR = TETAX(I,J-1)-AMIR
C
      IF (ALFAL.LE.ANGLE1.OR.ALFAL.GE.ANGLE2) GO TO 302
      XX(I,J) = XX(I-1,J)
      GO TO 307
302  IF(ALFAR.LE.ANGLE1.OR.ALFAR.GE.ANGLE2) GO TO 305
      XX(I,J) = XX(I,J-1)
      GO TO 307
305  XX(I,J) = (RX(I-1,J)-RX(I,J-1)+XX(I,J-1)*DTAN(ALFAR)+XX(I-1,J)*DTA
1N(ALFAL))/(DTAN(ALFAL)+DTAN(ALFAR))
C
307  IF (ALFAL.LE.0.000001.OR.ALFAL.GE.0.000001) GO TO 309
      RX(I,J) = RX(I-1,J)
      GO TO 315
309  IF (ALFAR.LE.0.000001.OR.ALFAR.GE.0.000001) GO TO 311
      RX(I,J) = RX(I,J-1)
      GO TO 315
311  RX(I,J) = RX(I,J-1)+(XX(I,J)-XX(I,J-1))*DTAN(ALFAR)
C
315  DKSI = DSQRT((RX(I,J)-RX(I,J-1))*2+(XX(I,J)-XX(I,J-1))*2)
      DETA = DSQRT((RX(I,J)-RX(I-1,J))*2+(XX(I,J)-XX(I-1,J))*2)
316  CONTINUE
C
      IF (RX(I,J).GE.0..AND.XX(I,J).GE.0.)GO TO 319
      KFIN = J-1
      WRITE (6,398)

```

```

AXS04330
AXS04340
AXS04350
AXS04360
AXS04370
AXS04380
AXS04390
AXS04400
AXS04410
AXS04420
AXS04430
AXS04440
AXS04450
AXS04460
AXS04470
AXS04480
AXS04490
AXS04500
AXS04510
AXS04520
AXS04530
AXS04540
AXS04550
AXS04560
AXS04570
AXS04580
AXS04590
AXS04600
AXS04610
AXS04620
AXS04630
AXS04640
AXS04650
AXS04660
AXS04670
AXS04680
AXS04690
AXS04700
AXS04710
AXS04720
AXS04730
AXS04740
AXS04750
AXS04760
AXS04770
AXS04780
AXS04790
AXS04800
AXS04810
AXS04820
AXS04830
AXS04840
AXS04850
AXS04860
AXS04870
AXS04880
AXS04890
AXS04900
AXS04910
AXS04920
AXS04930
AXS04940
AXS04950
AXS04960
AXS04970
AXS04980
AXS04990
AXS05000
AXS05010
AXS05020
AXS05030
AXS05040

```

```

398 FORMAT (' ','FURTHER POINTS ON THE CHARACTERISTICS ARE DIVERGENT')
C
C LOCATION OF THE NEW POINT HAS BEEN FOUND
C*****
C
C CALCULATE NOW P.M. ANGLE AND FLOW DIRECTION
319 APM = PMX(I,J-1)+PMX(I-1,J)-TETAX(I-1,J)+TETAX(I,J-1)
    IF(KD.EQ.2) GO TO 351
    BPM = DSIN(AMIR)*DSIN(TETAX(I,J-1))/RX(I,J-1)*DKSI
    CPM = DSIN(AMIL)*DSIN(TETAX(I-1,J))/RX(I-1,J)*DETA
351 PMX(I,J) = .5*(APM+BPM+CPM)
    APM = PMX(I,J-1)-PMX(I-1,J)+TETAX(I,J-1)+TETAX(I-1,J)
    TETAX(I,J) = .5*(APM+BPM+CPM)
C
C CALCULATE NOW THE MACH NUMBER FOR EACH POINT
C INITIAL GUESS AMX(I,J) = AMX(I-1,J)
    AMG = AMX(I-1,J)
    KZ = 0
    KH=0
    KL=0
354 IF (AMG.GT.2000.) GO TO 357
    D = DSQRT(AMG*AMG-1.)
    GO TO 358
357 D = AMG
358 IF (KZ.GE.50)GO TO 360
    KZ = KZ +1
    PMCAL = A2*DATAN(B2*D)-DATAN(D)
    DELNI = PMCAL-PMX(I,J)
    DEL = DABS(DELNI)
    IF (DEL.LT..000002) GO TO 360
    IF (DELNI.LT.0.)GO TO 356
    AMG = AMG*.98
    GO TO 354
356 IF (AMG.LT.5000.) GO TO 3560
    KFIN = J-1
    GO TO 399
3560 D1 = A3*AMG*AMG+1.
C*****
    DDELNI = DELNI
    IF (AMG.GT.20.) DDELNI = DELNI*(.95**KZ)
C*****
    AMG = AMG*(1.-DDELNI*D1/D)
    GO TO 354
360 AMX(I,J) = AMG
C
C CALCULATE NOW THE LOCAL TEMPERATURE, PRESSURE, VELOCITY, KNUDSEN
    C = AMX(I,J)*AMX(I,J)*A3+1.
    PRES = PSTG/(C*B1)
    TEMP = TSTG/C
    DN = PRES/(BOLTZ*TEMP)
C*****
    DENSF(I,J)=PRES/(RJ*TEMP)
    DDENS=DENSF(I-1,J-1)-DENSF(I,J)
    FP = .707/(DN*CXS)
    SCALE=DSQRT((XX(I,J)-XX(I-1,J-1))*2+(RX(I,J)-RX(I-1,J-1))*2)
    AKN=FP/SCALE
C
C
C*****
C    ALFALL = -ALFAL+PI
C    AF = ALFALL-PI/2.
C    AF = DABS(AF)
C    IF (AF.LT..000001) GO TO 370
C    BF = 1./DSQRT(1. + DTAN(ALFALL)**2)
C    SCALE=BF*(RX(I,J-1)-DTAN(ALFALL)*(XX(I,J-1)-XX(I-1,J))-RX(I-1,J))
C    SCALE = DABS(SCALE)
C*****
    SOUND = DSQRT(GAMA*RJ*TEMP)
    VEL = AMX(I,J)*SOUND
C
    COLF=4.*CXS*DN*DSQRT(BOLTZ*TEMP/(PI*GMM))
    P=VEL*DDENS/(SCALE*DENSF(I,J)*COLF)

```

C		AXS05770
C	PRINT RESULTS	AXS05780
	WRITE(6, 11)I, J, AMX(I, J), RX(I, J), XX(I, J), TETAX(I, J), TEMP, PRES, VEL,	AXS05790
	1AKN, FP, DN, P	AXS05800
399	CONTINUE	AXS05810
	STOP	AXS05820
	END	AXS05830
\$ENTRY		AXS05840

#### A.4 LIST OF MAIN SYMBOLS USED IN 'AXSYM'.

Parameter Name	Physical Name	Units	Type	Description
PAMB	ambient pressure	pascals	real	ambient atmosphere pressure
KD	K dimensions	-	integer	KD=2 for two dimensional jet KD=3 for axisymmetric ring jet
PI	$\pi$	-	real constant	
BOLTZ	Boltzmann constant	$\frac{\text{Joule}}{\text{degree}}$	real	$1.38032 \times 10^{-23}$ joules/degree
AVOG	Avogadro's constant	$\frac{\text{molecules}}{\text{mol}}$	real	$6.0225 \times 10^{26}$ 1/kmol
RG	Universal gas constant	$\frac{\text{Joule}}{\text{mol.deg}}$	real	8314.3
AM <sub>0</sub>	Mach No. (E.P)	-	real	Mach number at exit plane
TO	Temperature (E.P)	°k	real	Temperature at exist plane
PO	Pressure (E.P)	pascals	real	Pressure at exit plane
R1	Cylinder radius	m	real	Radius of the cylindrical vehicle
X1	0.5* nozzle width	m	real	Half width of nozzle
GAMA		-	real	averaged heat capacity ratio (jet)
DIAM	Molecule diameter	m	real	averaged molecular diameter (jet)
GM	Molecular mass	kg/kmol	real	averaged molecular weight (jet)
RJ	Gas constant	$\frac{\text{joules}}{\text{kJ.deg}}$	real	gas constant (jet)
CXS		m <sup>2</sup>	real	collision cross section (hard sphere)
GMM	mass of a molecule	kg	real	averaged mass of a molecule
N1			integer	number of divisions (characteristics from the exist plane)
N2			integer	number of characteristics in the Prandtl Meyer fan
A1			real	$\frac{\gamma - 1}{\gamma}$
B1			real	$\frac{\gamma}{\gamma - 1}$



## A.4 LIST OF MAIN SYMBOLS USED IN 'AXSYM' (CONTINUED)

Parameter Name	Physical Name	Units	Type	Description
A2			real	$[(\gamma + 1)/(\gamma - 1)]$
B2			real	$1/A2$
A3			real	$\frac{\gamma - 1}{2}$
C			real	$M^2 \frac{\gamma - 1}{2} + 1$
D,D1,D2			real	$M^2 - 1$
PSTG		pascals	real	stagnation pressure
TSTG		°k	real	stagnation temperature
DSTG		kg/m <sup>3</sup>	real	stagnation density
FSM			real	free stream Mach number
FST		°k	real	free stream temperature
FSD		kg/m <sup>3</sup>	real	free stream density
AMIT	$\mu_T$	radius	real	Mach angle (tail of P.M. fan)
ANIH	$\mu_H$	radius	real	Mach angle (head of P.M. fan)
PMT	$\nu_T$	radius	real	P.M. function (tail)
PMH	$\nu_H$	radius	real	P.M. function (head)
RAT,RAT2, E1			real	used to define a logarithmic division of the corner characteristics (50 lines in P.M. fan)(a linear division would have resulted in concentration of characteristics at high Mach numbers)
DELM			real	difference of Mach numbers
AM(I,J)	M		real 2-D array	Mach number at location (I,j)(used for the corner)
PM(I,J)	$\nu$	radius		P.M. function at location (I,j)(used for the corner)
AMI	$\mu$	radius	real	Mach angle
R(I,J)		m	real 2-D array	radius (or ordinate) at (I,j)
TETA(I,J)	$\theta$	radius	real 2-D array	flow direction at (I,j)

## A.4 LIST OF MAIN SYMBOLS USED IN 'AXSYM' (CONTINUED)

Parameter Name	Physical Name	Units	Type	Description
ALFAL		radians	real	angle of a left running characteristics
ALFAR		radians	real	angle of a right running characteristics
DENSF(I,J)		$\text{kg/m}^3$	real 2-D array	gas density at point IJ
AMCOR(I,J)			real 2-D array	Mach number at mesh points in region (1) (region 1 is bounded by the two tail characteristics)
TETAC(I,J)		radians	real 2-D array	flow direction at mesh points (region 1)
PMC(I,J)		radians	real 2-D array	P.M. function at mesh points (region 1)
XC(I,J)		m	real 2-D array	X coordinate at mesh points (region 1)
RC(I,J)		m	real 2-D array	radius (ordinate) at mesh points (region 1)
AMIL		radians	real	Mach angle far left running characteristics
AMIR		radians	real	Mach angle far right running characteristics
DKSI	$d\xi$	m	real	distance between mesh points
DETA	$d\eta$	m	real	distance between mesh points
DELNI	$dv$	radians	real	P.M. differential
AMG		radians		Mach number (used for iterative calculation)
PRES				pressure
TEMP				temperature
DN				number density
FP				mean free path
AKN				Knudsen number
SOUND				speed of sound
VEL				absolute local velocity
DDENS				density difference between two points along a streamline
COLF				collision frequency
P				breakdown parameter



## A.5 AXSYM PROGRAM USER'S GUIDE

### 1. Input data:

ambient pressure	PAMB
Mach number (Exit plane)	AMO
Temperature (Exit plane)	TO
Pressure (Exit plane)	PO
Half width of nozzle	X1
Radius of nozzle ring	R1
Specific heat ratio of jet gas	GAM1A
average molecular diameter (jet)	DIAM
average molecular weight	GM

### 2. Options for flow geometry

two dimensional flow	KD=2
ring jet	KD=3
default condition	KD=3

### 3. Resolution of mesh points

to change the resolution of the mesh points

a - change N1 and N2 as necessary

b - change 'DIMENSIONS' according to new values of  $N_1$  and  $N_2$

c - define distribution of Mach lines in the Prandtl-Meyer fan as required  
(program lines 126-131)

### 4. Execution commands:

After copying program into USER'S FILE:

```
WATFIV AXSYM * (XTYPE
```

The program will run on user's terminal under WATFIV. A soft copy of program listing and output listing will be stored in user's disk named AXSYM LISTING.

5. Hard Copy

PRINT AXSYM LISTING.

6. Program Outputs.

All necessary outputs are automatically listed by the program.

Figure (20) and Figure (21) show the resulting mesh of characteristics calculated for an altitude of 200 km for  $M_0=4$  and  $M_0=2$  respectively.

In these figures we also show some isotherms and the limit where the breakdown parameter equals 0.05. These lines are plotted (manually) using interpolation procedures. Data along the breakdown line is input data for the molecular flow.

B.1 DATA ORGANIZATION

Because of the large number of molecules, cells, regions and sectors in the simulation and the large number of data related to each molecule, each cell and region to be stored, special precautions should be taken in order not to overflow the available computer memory.

The following data organization was used in SIMUL. Figure 22 shows the geometry of one sector.

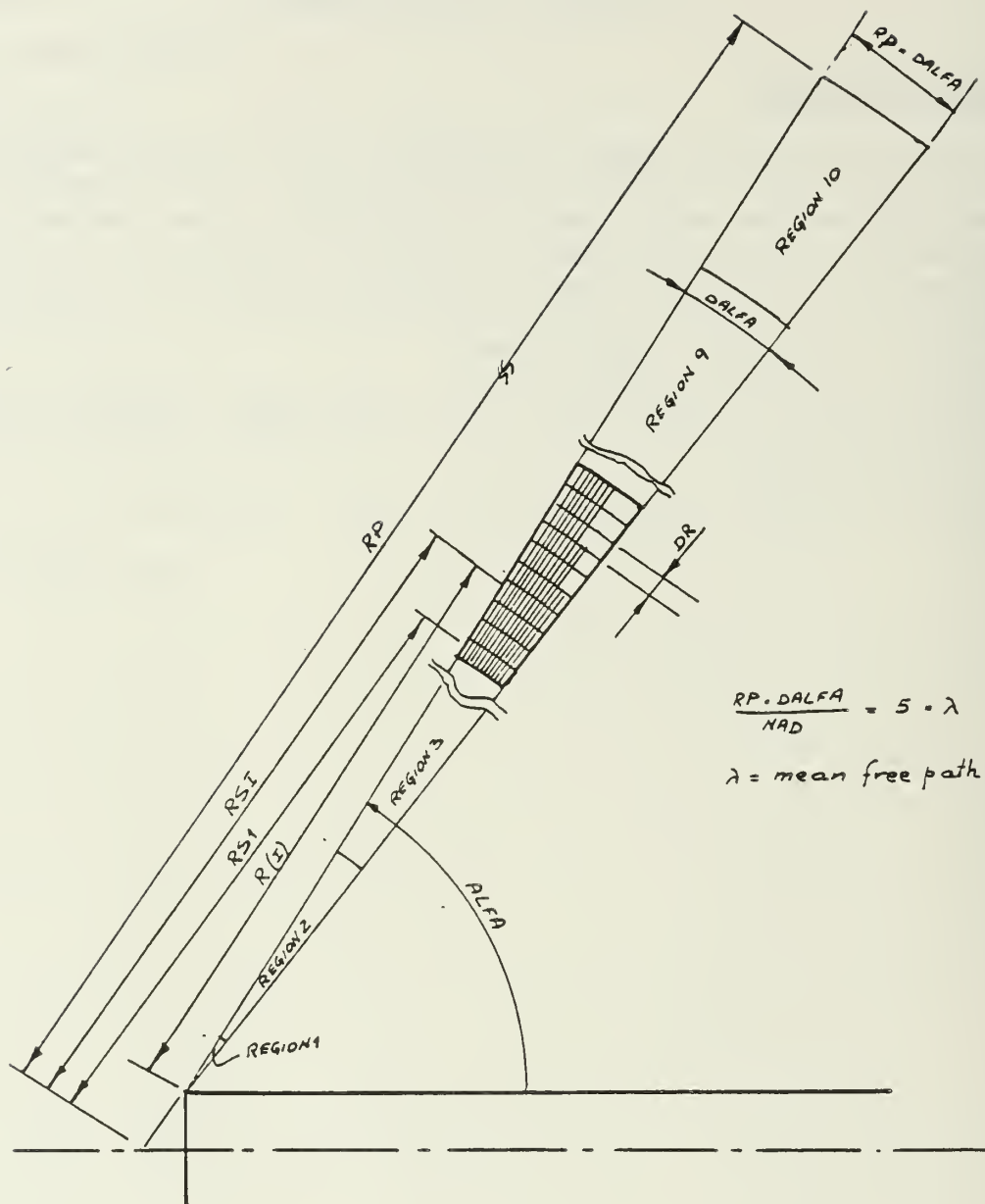


Figure 22. Definition of sector geometry and cell volume.

$$\text{cell volume} = v(I) = R(I) * DDALFA * DR * RSI * DFI.$$

$$\frac{v(I)}{v(1)} = \frac{R(I) \cdot RSI(I)}{R(1) \cdot RSI}$$

$v(1)$  is the volume of smallest cell in a region.

1. Tables of Molecules and Their Parameters

P1(L,N1M) - Light molecules of the jet

P2(L,N2M) - Heavy molecules of the jet

P3(L,N3M) - Ambient gas molecules (not used in the present program)

N1M, N2M, N3M - maximum number of molecules in simulation. Number of active molecules may be smaller or equal to  $(N \leq I \leq M)$ .

L=1,2,3 - cartesian components of velocity  $v_x, v_y, v_z$  [m/s]

L=4 - radial coordinate [meters]

if  $r=-99$  the particular molecule is inactive

L=5 - angular coordinate [radians].

This table is generated each time the simulation is initiated for a region.

That means, the same group of molecules (as stored in the computer memory) is used to simulate the flow in all regions in the computation domain.

2. C(M,I,j) Table of Cells (in a Region)-Real Data

- I = 1,10 - radial index of the cell in a region
- j = 1,10 - angular index of the cell in a region
- M = 19 - radial coordinate of cell center
- M = 20 - angular coordinate of cell center
- M = 1,9 - time parameter for collisions of different species in a  
cell
- M = 10-18 - maximum relative velocity expected for collisions of  
different species in a cell



3. IP(N1A+N2A+N3A) Table of the addresses of the active molecules  
arranged in order of their species and in the order of their cells

IC(N,I,j) - table of cells (in a region) integer data

I = 1,10 - radial index

j = 1,10 - angular index

N = 1 - number of molecules (spec 1)

N = 2 - number of molecules (spec 2)

N = 3 - number of molecules (spec 3)

N = 4 - (starting address - 1) of molecules as ordered in (IP).

4. Reg(N,kR,kS) Data table for a specific region (real)

kR = 1,10 - index of a region in a sector

kS = 1,20 - index of the sector

N = 1 - Df1 = differential angle (axisymmetric)

DF1 is a weighting factor

N = 2 - DN1 = number density (species 1)

N = 3 - DN2 = number density (species 1)

N = 4 - VOL 1 = actual volume of smallest cell in kR

N = 5 - AREA1 INPUT area of smallest cell in kR

## 5. Region Geometry and Input Flux

R(10) - polar radius of a cell in a region

A(10) -  $\frac{\text{input area of a cell}}{\text{input area of smallest cell}}$

VOL(10) -  $\frac{\text{volume of a cell}}{\text{volume of smallest cell}}$

M1(10) - initial number of molecules in cells

(equal number of molecules of either jet species)

F1(10), F2(10) - input flux through high pressure starting line

(spec 1), (spec 2)

## 6. Input Flux From High Pressure Starting Line

FWP1(N,I,j)

FWP2(N,I,j)

↑↑↑↑  
| species  
| positive (input molecules)  
| west  
| flux

I = 1,10 - number of the cell along the starting line in a region  
(j)

j = 1,10 - number of the region along the starting line

N = 1 - molecular flow for a given DF1

N = 2 - mean molecular velocity Vx

N = 3 - mean molecular velocity Vy

N = 4 - mean input gas temperature

## 7. Output Flux

FNN1(4,NAD,kR)

FSN1(4,NAD,kR)

FNN2(4,NAD,kR)

FSN2(4,NAD,kR)

↑↑  
| negative (output)  
| north

↑↑  
| negative (output)  
| south

kR - number of region in the sector

NAD - angular location

The first index include the same parameters as (FWP1)

FEN1(4,NRD)

FWN1(4,NRD)

FEN2(4,NRD)

FWN2(4,NRD)

↑↑  
| negative  
| east

↑↑  
| negative  
| west

NRD → radial location

FEN1, FEN2, FWN1, FWN2 are necessary for iterations within one sector.

#### 8. Sampling of Output Flux from a Sector

After averaging they are transferred to FOE1, FOE2, FOW1, FOW2.

FOW1(4, kC, kR, kS)

output flow to the west

FOW2(4, kC, kR, kS)

FOE1(4, kC, kR, kS)

output flow to the east

FOE2(4, kC, kR, kS)

kC - radial location (cell) of flow in a region

kR - index of a region in a sector

kS - index of the sector

The first index (4 parameters) include the same parameters as FEPl( ).



## B.2 MOLECULAR SIMULATION FOR A GIVEN REGION

After we define the geometry of the whole sector resulting from the region geometry and cell geometry, we may start with the molecular simulation. This includes:

- initial setting of molecules in cells
- molecules are moved according to the time increment  $\Delta t$
- new molecules are generated according to input (or output) flows
- collisions calculations
- integration of flow parameters for average parameters calculations
- repetition of the whole procedure as long as necessary to obtain reasonable statistical averaging
- calculate averages and flow weighting.

These routines are the core of the program and must be repeated for all regions in a sector and (or all sectors in the domain where the collisions are significant).

In the following sections we bring a detailed description of this part of the program.

### 1. Initial Setting of Molecules in Cells

- a. The initial number of available molecules in a simulation is larger than the number of active molecules ( $P1(\text{data}, \text{number of molecules})$ ,  $P2(\text{data}, \text{number of molecules})$  are the vectors used for species 1 and 2)
- b. an inactive molecule is defined as

$$[P1(4,N) \text{ or } P2(4,N)] = -99$$

- c. calculation of number of molecules to be set in each cell
- d. calculation of cell coordinates
- e. deactivation of all available molecules in simulation
- f. definition of molecules coordinates

P1(4,N), P2(4,M) are polar radiuses

P1(5,N), P2(5,M) are angular coordinate in radians

All molecules in a cell are set at random locations within the cell.

- g. definition of molecular velocities

P1(1,N), P2(1,M) velocity in X direction

P1(2,N), P2(2,M) velocity in Y direction

P1(3,N), P2(3,M) velocity in Z direction

Thermal velocities are random function of temperatures and are added to the mean velocity as defined at initial boundary  $ALFA_0$  of the region.

As the thermal velocity has a Boltzmann distribution the thermal velocity setting is based on rejection-acceptance methods (for more details see Bird [4] Appendix D).

- h. reset collision timers and relative velocity

i. reset general time counter: Time = 0

### 3. The Simulation

- a. Move all molecules according to their velocity ( $V_x$ ,  $V_y$ ) and find their new coordinates

Note - A routine designed to calculate the collisions with the wall was included in the program; if the region (or sector) is bounded by the solid wall, the collision is calculated - resulting new velocities and directions and counted for wall flux calculations. If the program is stopped at an angle where the flow becomes collisionless, this routine becomes irrelevant and other type of calculations should be designed.

- b. Output flow counting: all molecules that leave the region are counted and stored in specific vectors which are used as inputs to other regions. The output vectors are (X represent 1 or 2 for the two species in the program (FSNX))

FSNX(1,j,kR) → "south" boundary

FNNX(1,j,kR) → "north" boundary

FENX(1,I) → "east" boundary

FWNX(1,I) → "west" boundary

I represents the radial location index of the cell

j represents the angular location index

kR is the index of the region within a sector

Note - all molecules that move to

(j=NAD+1)(I=NRD+1) are placed in FENX(1,NRD)

(j= -1)(X=NRD+1) are placed in FWNX(1,NRD)

(j=NAD+1)(I= -1) are placed in FSNX(1,1,kR)

(j= -1)(I= -1) are placed in FWNX(1,1,kR)

This was done only for simplification reasons.

- c. Generation of new molecules due to input flows. Through the four sides E,N,W,S, of a region, molecules are allowed to enter the region according with the flows coming from the neighbouring regions:

for "W" side of the region in sector 1

FWPX(P,I,kR)

for other sectors

FOEX(P,I,kR,kS-1)

for "E" side of any region

FOWX(P,I,kR,kS+1)

for "N" side of any region

FSNX(P,j,kR+1)

for "S" side of the region

FNNX(P,j,kR-1)

The first parameter of all these arrays represent:

P = 1 - number flux (real number)

P = 2 - velocity component - ( $V_x$ )

P = 3 - velocity component - ( $V_y$ )

P = 4 - gas temperature

Note 1 - because every region has a different size of angle DFI, all fluxes have to be adjusted accordingly.

Note 2 - input fluxes are calculated, adjusted and stored as real numbers. The number of input molecules are by definition integers. In order not to "lose" molecules, the number of input molecules is increased by 1 on a random basis. (The average of many runs will result in the accurate average input flow.)

New molecules are set at random locations on the boundary of the specific cells and at random time within DTM. Then each molecule is allowed to enter the region according to its initial coordinate and velocity. At the end of the time interval the new location and velocity is stored in molecule array P1 or P2. If DTM is chosen to be too large and cell size is small (total region size too small) some molecules may cross the region and will not be counted in the simulation of the specific region. In order not to "lose" molecules:

- (a) DTM should be decreased
- (b) count these molecules as output fluxes from the specific region. (This is recommended only if there is no other choice.)

Note - Because the arrays of input flows store only averaged data for the molecules, the thermal velocity of each new molecule is calculated according to the Boltzmann distribution as a function of the averaged temperature.

d. Rearrangement of molecules in cells. Before collisions are calculated all simulated molecules which have been let to move and generated have to be rearranged and recounted for each cell.



The array  $IC(k,I,j)$  contains integer data for each cell

$(I,j)$

$k = 1$  - is the number of molecules of species 1

$k = 2$  - is the number of molecules of species 2

$k = 3$  - is the number of molecules of species 3 (not used)

$k = 4$  - is the (address-1) of the first molecule in the cell

related to the vector  $IP(M)$

Vector  $IP(M)$  contains the list of the simulated molecules arranged in the order of species in cells and cells respectively. The following is a graphic description of  $IP(M)$

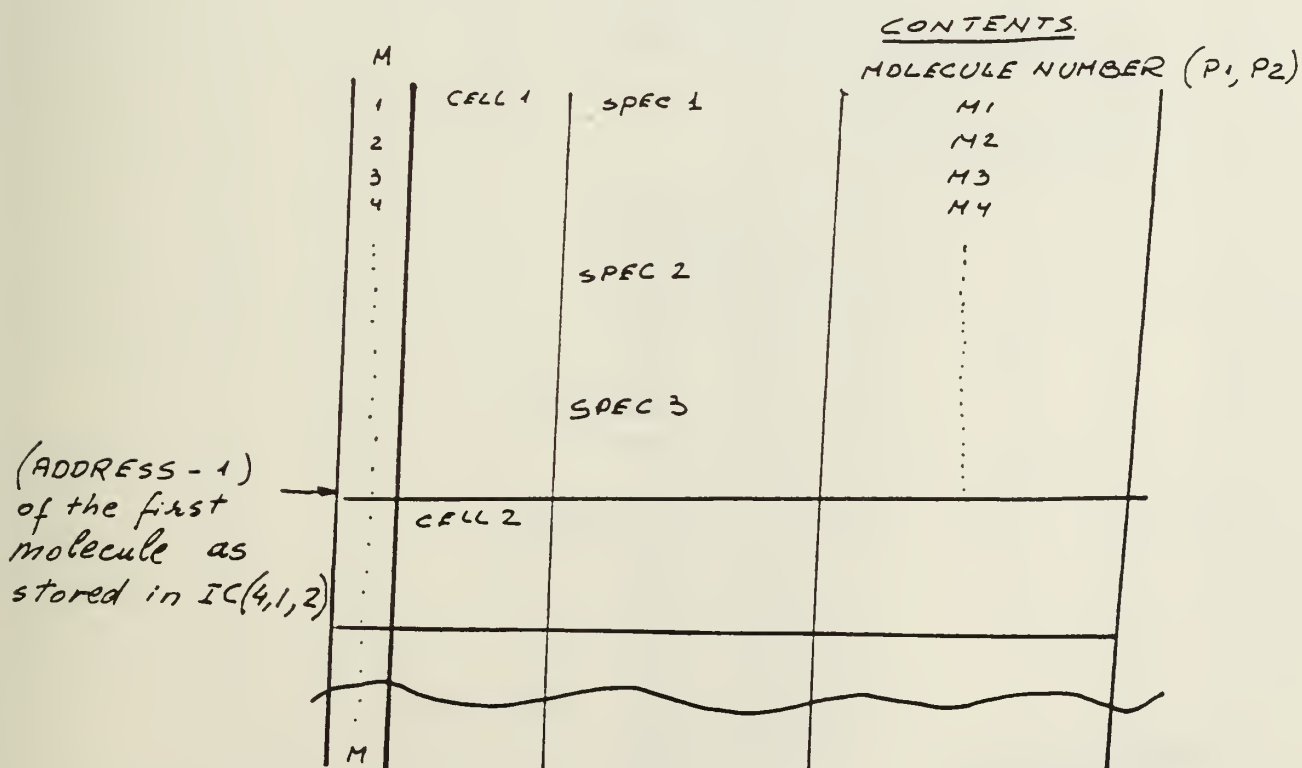


Figure 23. Vector  $IP(M)$

### B.3 SIMUL Program Flowchart

A simplified flowchart for the Monte Carlo simulation of the molecular flow is shown in Figure 24. The program is designed to solve the ring axisymmetric jet flow, however, minor changes may be done to enable a different geometry.



## B.4 SIMUL Program Listing

```

C      PROGRAM SIMUL
C THIS PROGRAM IS DESIGNED TO CALCULATE THE MOLECULAR FLOW OUTSIDE THE
C CONTINUUM REGION FLOW OF A HIGHLY UNDEREXPANDED AXISYMMETRIC RING JET.
C RESULTS FOR THE CONTINUUM FLOW MAY BE OBTAINED FROM 'AXSYM' PROGRAM
C WHICH GIVES THE 'METHOD OF CHARACTERISTICS' ISENTROPIC SOLUTION.
C THE BOUNDARY BETWEEN THE CONTINUUM AND MOLECULAR FLOW IS DEFINED BY
C THE BREAKDOWN PARAMETER 'P' AS PROPOSED BY G. A. BIRD.
C THE 'MOLECULAR DOMAIN' IS DIVIDED INTO POLAR DIVISIONS MAKING A SET
C OF SECTORS. EACH SECTOR IS SUBDIVIDED INTO 10 REGIONS
C NO APRIORY INFORMATION ABOUT THE GEOMETRY OF THE DIFFERENT
C REGIONS IS AVAILABLE, THEREFORE, MANUAL INTERVENTION MAY BE
C REQUIRED WHEN MOVING FROM ONE SECTOR TO THE OTHER. AN ACCEPTABLE
C GEOMETRY WILL RESULT A REASONABLE NUMBER OF SIMULATED MOLECULES.
C EACH REGION IS SUBDIVIDED INTO A NUMBER OF CELLS WITH A GEOMETRY
C DEFINED BY A POLAR MESH. A NUMBER OF MOLECULES IS SET IN EACH CELL
C PROPORTIONAL TO THE CELL VOLUME. THE BOUNDARY CONDITIONS FOR EACH
C REGION REQUIRE INPUT AND OUTPUT FLUX OF MOLECULES. NO APRIORY
C INFORMATION ON THE FLUX IS AVAILABLE. IT WILL BE CALCULATED IN
C AN ITERATIVE MODE.
C IF THE BOUNDARY CONDITIONS FOR ALL CELLS ARE CONSTANT THE NUMBER
C FLUX INTO EACH CELL IS PROPORTIONAL TO CELL WALL AREA.
C TO DECREASE THE ERROR WHEN INTRODUCING MOLECULES -(INTEGER NUMBER)-
C DUE TO THE INPUT FLUX -(REAL NUMBER), AN ADDITIONAL MOLECULE IS
C GENERATED ON THE BASIS OF RANDOM NUMBERS SUCH THAT THE AVERAGE OF A
C LARGE NUMBER OF SAMPLINGS WILL EQUAL THE REAL INPUT FLUX.
C*****
C      THE MONTE CARLO SIMULATION
C THE JET GAS IS COMPOSED OF TWO SPECIES OF MOLECULES
C AMBIENT GAS IS REGARDED AS ONE SPECIES
C THE MOLECULAR MODEL IS - 'HARD SPHERE MOLECULE'
C NETWORK DEFINITION
C THE MAXIMUM RADIUS (POLAR) OF THE DOMAIN IS ASSUMED TO BE RP=15 M.
C THE ANGLE OF THE BOUNDARY BETWEEN CONTINUUM AND MOLECULAR FLOW
C AND THE SOLID WALL IS
C      ALFA (CALCULATED IN PROGRAM 'AXSYM')
C DEFINE A POLAR SECTOR WITH A RADIUS 'RP' AND AN ANGLE OF
C      DALFA =5*MFP*NAD/RP
C THIS SECTOR IS SUBDIVIDED INTO A NUMBER OF RADIAL DIVISIONS
C MAKING 'N' REGIONS FOR SIMULATION CALCULATIONS FOR EACH 'DALFA'.
C EACH REGION IS DEVIDED INTO
C      NAD -ANGULAR DIVISIONS (15) WITH AN ANGLE OF DDALFA=5*MFP/RP
C      NRD -RADIAL DIVISIONS (10)
C      MAKING NAD*NRD CELLS
C THE SMALLEST CELL CONTAINS 'MIN' MOLECULES
C      MIN = 15
C TOTAL NUMBER OF ACTIVE MOLECULES IN A REGION IS LIMITED TO 6000
C (3000 MOLECULES OF EACH SPECIES.)
C THE WIDTH OF A CELL DEFINED BY ANGLE 'DFI' MAY NOW BE EVALUATED.
C (MIN/NUMBER DENSITY = ACTUAL CELL VOLUME)
C*****
C      DEFINE PARAMETERS.....BLOCK 1
C      COMMON IX
C      DIMENSION P1(5,3000),P2(5,3000),IP(6000),C(20,10,15),
C      *IC(4,10,15),REG(5,10,20)
C      DIMENSION DFI(10),DN(10)
C      DIMENSION R(10),A(10),VOL(10),M1(10),F1(10),F2(10)
C      DIMENSION SS(2,10,10)
C      DIMENSION FEN1(4,10),FWN1(4,10),FNN1(4,15,10),FSN1(4,15,10)
C      DIMENSION FEN2(4,10),FWN2(4,10),FNN2(4,15,10),FSN2(4,15,10)
C      DIMENSION FWp1(4,10,10),FWP2(4,10,10)
C      DIMENSION FOE1(4,10,10,20),FOW1(4,10,10,20)
C      DIMENSION FOE2(4,10,10,20),FOW2(4,10,10,20)
C      DIMENSION NM(3),VRC(3),TOC(3,3),SPEC(3,5),NCOL(10,15)
C P1,P2,P3 CONTAIN INFORMATION ON UP TO M SIMULATED MOLECULES
C P1(1,N),P1(2,N),P1(3,N) ARE U,V,W VELOCITY COMPONENTS (CARTESIAN)
C P1(4,N),P1(5,N) ARE R AND TETA COORDINATES (POLAR)
C
C IP(M) ARE THE M MOLECULES ARRANGED IN THE ORDER OF THEIR CELLS
C C(20,I,J) CONTAINS INFORMATION ON UP TO I*J CELLS
C*****
C ALFA IS THE ANGLE WHERE THE BREAKDOWN PARAMETER EQUALS .05

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C FPM IS THE MEAN FREE PATH AT ALFA. SIM00730
C***** SIM00740
C SET GENERAL CONSTANTS.....BLOCK 2 SIM00750
  IX = 529814367 SIM00760
  PI = 3.141593 SIM00770
  BOLTZ = 1.38044E-23 SIM00780
  P0 = 101325. SIM00790
  T0 = 273. SIM00800
  AVOG = 2.68699E+25 SIM00810
  RG = 8314. SIM00820
C***** SIM00830
C SET PROGRAM CONSTANTS.....BLOCK 3 SIM00840
  ISPEC=2 SIM00850
  R1 = 2.5 SIM00860
  DR=.15 SIM00870
  RP=15. SIM00880
C NUMBER OF SIMULATED MOLECULES IN SMALLEST CELL (DIFFERENT SPECIES) SIM00890
  NMOL1=3000 SIM00900
  NMOL2=3000 SIM00910
  NMOL3=0 SIM00920
  MIN = 15 SIM00930
  N1M = 15 SIM00940
  N2M = 15 SIM00950
C NUMBER OF DIVISIONS IN A SIMULATED REGION SIM00960
  NAD = 15 SIM00970
  NRDS=10 SIM00980
  NRD=10 SIM00990
  NRD1= 3 SIM01000
  NIS =2 SIM01010
C R1 IS THE RADIUS OF THE CYLINDER (WALL) SIM01020
C R(I) IS THE RADIAL COORDINATE OF CELL(I) SIM01030
C VOL(I) IS THE RATIO BETWEEN VOLUMES OF CELL(I) AND CELL(1) SIM01040
C A(I) IS THE RATIO BETWEEN INPUT FLOW AREAS OF CELL(I) AND CELL(1) SIM01050
C DR IS THE RADIAL SIZE OF A CELL (RADIAL INCREMENT) SIM01060
C INPUT THE FOLLOWING AS 'DATA' OR 'READ' STATEMENTS ***** SIM01070
  TETA = 1.2 SIM01080
C TETA IS THE FLOW DIRECTION (RADIAN) ON THE BREAKDOWN LINE AS FOUND SIM01090
C FROM THE AXSYM PROGRAM. V0 IS THE FLOW VELOCITY (M/SEC) SIM01100
  V0 = 2100 SIM01110
  VOX = V0*COS(TETA) SIM01120
  VOY = V0*SIN(TETA) SIM01130
  TWALL=300. SIM01140
  DN01=1.1E21 SIM01150
  DN02=1.1E21 SIM01160
  DTM=1.5E-6 SIM01170
C SET DATA FOR THE DIFFERENT SPECIES SIM01180
C MOLECULAR MASS SIM01190
  SPEC(1,1)=4./AVOG SIM01200
  SPEC(2,1)=40./AVOG SIM01210
  SPEC(3,1)=29./AVOG SIM01220
C MOLECULAR DIAMETER SIM01230
  SPEC(1,2)=2.19E-10 SIM01240
  SPEC(2,2)=4.00E-10 SIM01250
  SPEC(3,2)= SIM01260
  SPEC(1,3)= SIM01270
  SPEC(2,3)= ADDITIONAL DATA IF REQUIRED SIM01280
  SPEC(3,3)= SIM01290
C***** SIM01300
C THERMAL VELOCITIES AT WALL TEMPERATURE SIM01310
  VWM1=SQRT(2.*BOLTZ*TWALL/SPEC(1,1)) SIM01320
  VWM2=SQRT(2.*BOLTZ*TWALL/SPEC(2,1)) SIM01330
C SIM01340
C***** SIM01350
C INITIALISATION..RESET ALL SAMPLING VARIABLES SIM01360
  DO 80 I=1,NRD SIM01370
  DO 80 JR=1,10 SIM01380
  DO 80 JS=1,20 SIM01390
  DO 80 KPAR=1,4 SIM01400
  FOE1(KPAR,I,JR,JS)=0. SIM01410
  FOE2(KPAR,I,JR,JS)=0. SIM01420
  FOW1(KPAR,I,JR,JS)=0. SIM01430
  80 FOW2(KPAR,I,JR,JS)=0. SIM01440

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C*****SIM01450
C SET INPUT PARAMETERS.....BLOCK 3 SIM01460
  ITER=1 SIM01470
C RETURN TO 3000 FOR ADDITIONAL ITERATIONS SIM01480
  3000 KS=1 SIM01490
    ALFA=1.3 SIM01500
    FPM=.001 SIM01510
    TEMP=40. SIM01520
    VTER1=SQRT(2.*BOLTZ*TEMP/SPEC(1,1)) SIM01530
    VTER2=SQRT(2.*BOLTZ*TEMP/SPEC(2,1)) SIM01540
    IF (ITER.GT.1)GO TO 2005 SIM01550
    DO 2001 I=1,10 SIM01560
      REG(2,I,1)=DN01 SIM01570
      REG(3,I,1)=DN02 SIM01580
    DO 2001 J=1,10 SIM01590
      FWP1(2,I,J)=V0X SIM01600
      FWP2(2,I,J)=V0X SIM01610
      FWP1(3,I,J)=V0Y SIM01620
      FWP2(3,I,J)=V0Y SIM01630
      FWP1(4,I,J)=TEMP SIM01640
    2001 FWP2(4,I,J)=TEMP SIM01650
  2005 CONTINUE SIM01660
C SIM01670
C*****SIM01680
C*****SIM01690
C*****SIM01700
C DEFINE SECTOR GEOMETRY.....BLOCK 4 SIM01710
C KR IS THE INDEX FOR A REGION IN ONE SECTOR SIM01720
C RETURN TO 1000 FOR NEXT SECTOR SIM01730
  DALFA=0. SIM01740
  1000 KR=1 SIM01750
C RESET SAMPLING OF FLOW VARIABLES (N AND S) SIM01760
  DO 81 I=1,NAD SIM01770
    DO 81 JR=1,10 SIM01780
      DO 81 KPAR=1,4 SIM01790
        FNN1(KPAR,I,JR)=0. SIM01800
        FNN2(KPAR,I,JR)=0. SIM01810
        FSN1(KPAR,I,JR)=0. SIM01820
      81 FSN2(KPAR,I,JR)=0. SIM01830
C ALFA IS THE STARTING ANGLE OF THE PRESENT SECTOR. PREVIOUS DALFA SIM01840
C SUBTRACTED FROM PREVIOUS ALFA (PREVIOUS SECTOR) GIVES THE NEW ALFA. SIM01850
  ALFA=ALFA-DALFA SIM01860
  DALFA = 5.*FPM*FLOAT(NAD)/RP SIM01870
C DALFA IS THE ANGLE OF THE SECTOR SIM01880
  IF(DALFA.GT.ALFA/2.)DALFA=ALFA/2. SIM01890
  IF(DALFA.GT.(FPM*75/RP))DALFA=ALFA SIM01900
  IF(ALFA.LE.0.)GO TO 2000 SIM01910
  DDALFA = DALFA/FLOAT(NAD) SIM01920
C DDALFA IS THE ANGLE OF A SINGLE CELL IN THE SECTOR SIM01930
  ALFAJ=ALFA-DALFA/2. SIM01940
C SIM01950
C*****SIM01960
C DEFINE REGION KR IN SECTOR.....BLOCK 5 SIM01970
C RETURN TO 2000 FOR THE NEXT REGION KR SIM01980
  2000 CONTINUE SIM01990
C RESET SAMPLING OF FLOW VARIABLES PER REGION SIM02000
  DO 82 I=1,NRD SIM02010
    DO 82 KPAR=1,4 SIM02020
      FEN1(KPAR,I)=0. SIM02030
      FEN2(KPAR,I)=0. SIM02040
      FWN1(KPAR,I)=0. SIM02050
    82 FWN2(KPAR,I)=0. SIM02060
  MT=0 SIM02070
  NRD=NRDS SIM02080
  IF(KR.LT.2)NRD=NRD1 SIM02090
  DO 100 I = 1,NRD SIM02100
C POLAR RADIUS MEASURED FROM THE NOZZLE LIP SIM02110
  R(I)=(FLOAT(I)-.5)*DR SIM02120
  IF(KR.EQ.2)R(I)=R(I)+FLOAT(NRD1)*DR SIM02130
  IF(KR.GE.3)R(I)=R(I)+(FLOAT(NRD1+(KR-2)*NRD))*DR SIM02140
C RSI IS THE RADIUS MEASURED FROM THE AXIS OF SYMMETRY SIM02150
  RSI=R(I)+R1/SIN(ALFAJ) SIM02160

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      IF(I.EQ.1)RS1=RSI
      A(I) = RSI/RS1
      VOL(I) = R(I)*RSI/(R(1)*RS1)
      M1(I) = MIN*VOL(I)
      MT = MT+M1(I)
C M1() IS THE INITIAL NUMBER OF MOLECULES IN EACH CELL
      DN1=REG(2,KR,KS)
      DN2=REG(3,KR,KS)
      REG(1,KR,KS)=FLOAT(MIN)/((DN1*R(1)*RS1*DR*DDALFA)
C DFI (REG(1,KR,KS)) IS THE SPHERICAL (AXISYMMETRIC) ANGLE
C THIS ANGLE HAS DIFFERENT VALUES FOR EACH KR, THEREFORE IT IS A
C WEIGHTING FACTOR.+++++
C (FNN1,FNN2,FSN1,FSN2,FEN1,FEN2,FWN1,FWN2) * DFI
C (FOE1,FOE2,FOW1,FOW2) * DFI
      DA1 = RS1*DR*REG(1,KR,KS)
C DA1 IS THE INPUT AREA OF CELL(1)
      FN1 = V0*DA1*REG(2,KR,KS)*SIN(ALFA-TETA)
      FN2 = V0*DA1*REG(3,KR,KS)*SIN(ALFA-TETA)
C TETA IS THE ANGLE BETWEEN FLOW DIRECTION AND THE WALL
      F1(I) = FN1*A(I)
      F2(I) = FN2*A(I)
C
      99 FWP1(1,I,KR)=F1(I)
      100 FWP2(1,I,KR)=F2(I)
C .....
C DEFINE ACTUAL VOLUME AND INPUT AREA OF SMALLEST CELL IN REGION
      REG(4,KR,KS)=R(1)*DDALFA*DR*RS1*REG(1,KR,KS)
      REG(5,KR,KS)=DA1
C FN1 IS THE NUMBER FLUX TO THE SMALLEST CELL (REAL NUMBER) PER SECOND.
C INPUT NUMBER OF MOLECULES (INTEGER) WILL BE INTEGRATED TO MAKE AN
C AVERAGE OF FN1.
C F1(I) IS THE INPUT FLUX TO CELL(I)
C FOLLOWING ARE THE REGION BOUNDARIES
      102 RMIN=R(1)-.5*DR
      RMAX=RMIN+DR*FLOAT(NRD)
      TMAX=ALFA
      TMIN=ALFA-DALFA
      DO 9102 I = 1,NRD
      WRITE (6,9101)R(I),A(I),VOL(I),M1(I),F1(I)
9101  FORMAT(' ',3F13.5,I10,E15.5)
9102  CONTINUE
      MTT= MT*NAD
      WRITE(6,103)MT,MTT
      103 FORMAT (' INITIAL NUMBER OF SIMULATED MOLECULES IS= ',I5,' PER SPES
      XCIES PER DDALFA, TOTAL NUMBER IS', I5)
C*****
C DEFINE INPUT FLOWS TO KR W,E,N,S.....BLOCK 6
C
C
C*****
C CALCULATE INITIAL NUMBER OF MOLECULES IN CELLS.....BLOCK 7
C SET INITIAL STATE OF GAS..... '' 8
C
      DO 150 I =1,NRD
      DO 150 J= 1,NAD
C
C SET SIMULATED MOLECULES IN THEIR CELLS
      IC(1,I,J) = M1(I)
      IC(2,I,J) = M1(I)
      IC(3,I,J) = 0
C
C SET CELL COORDINATES
      C(19,I,J) = R(I)
      C(20,I,J) = ALFA-(FLOAT(J)-.5)*DDALFA
      150 CONTINUE
C*****
C DEACTIVATE ALL MOLECULES
      DO 170 N = 1,NMOL1
      P1(4,N) = -99.
      170 P2(4,N) = -99.

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SIM02170
SIM02180
SIM02190
SIM02200
SIM02210
SIM02220
SIM02230
SIM02240
SIM02250
SIM02260
SIM02270
SIM02280
SIM02290
SIM02300
SIM02310
SIM02320
SIM02330
SIM02340
SIM02350
SIM02360
SIM02370
SIM02380
SIM02390
SIM02400
SIM02410
SIM02420
SIM02430
SIM02440
SIM02450
SIM02460
SIM02470
SIM02480
SIM02490
SIM02500
SIM02510
SIM02520
SIM02530
SIM02540
SIM02550
SIM02560
SIM02570
SIM02580
SIM02590
SIM02600
SIM02610
SIM02620
SIM02630
SIM02640
SIM02650
SIM02660
SIM02670
SIM02680
SIM02690
SIM02700
SIM02710
SIM02720
SIM02730
SIM02740
SIM02750
SIM02760
SIM02770
SIM02780
SIM02790
SIM02800
SIM02810
SIM02820
SIM02830
SIM02840
SIM02850
SIM02860
SIM02870
SIM02880

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C
C*****
C SET INITIAL STATE OF THE GAS (LOCATION AND VELOCITY OF MOLECULES)
C
    NADR1 = 0
    NADR2 = 0
    DO 200 I = 1,NRD
    DO 200 J = 1,NAD
    NM1 = IC(1,I,J)
C
    DO 205 N = 1,NM1
    NADR1 = NADR1 + 1
    CALL RANDU(PP)
    P1(4,NADR1) = C(19,I,J)+DR*(PP-.5)
    CALL RANDU(P)
    P1(5,NADR1) = C(20,I,J)+DDALFA*(P-.5)
C
    DO 205 NV = 1,3
203 CALL RANDU(P)
    V = -3.+6.*P
    B = EXP(-V*V)
    CALL RANDU(P)
    IF(B.LT.P) GO TO 203
    P1(NV,NADR1) = P*SIN(B)*VTER1
    IF(NV.EQ.1)P1(NV,NADR1)=P1(NV,NADR1)+VOX
    IF(NV.EQ.2)P1(NV,NADR1)=P1(NV,NADR1)+VOY
205 CONTINUE
C
C REPEAT PROCEDURE FOR SPECIES 2
    NM2 =IC(2,I,J)
    DO 210 N = 1,NM2
    NADR2 = NADR2+1
    CALL RANDU(P)
    P2(4,NADR2) = C(19,I,J)+DR*(P-.5)
    CALL RANDU(P)
    P2(5,NADR2) = C(20,I,J)+DDALFA*(P-.5)
C
    DO 210 NV = 1,3
207 CALL RANDU(P)
    V = -3.+6.*P
    B = EXP(-V*V)
    CALL RANDU(P)
    IF(B.LT.P) GO TO 207
    P2(NV,NADR2) = P*SIN(B)*VTER2
    IF(NV.EQ.1)P2(NV,NADR2)=P2(NV,NADR2)+VOX
    IF(NV.EQ.2)P2(NV,NADR2)=P2(NV,NADR2)+VOY
210 CONTINUE
C
C WHEN NECESSARY, REPEAT PROCEDURE FOR SPECIES 3.
200 CONTINUE
C
C*****
C DEFINE HERE ALL COLLISION PARAMETERS TO BE INCLUDED IN SIMULATION
C
C RESET COLLISION TIMERS
    DO 19 I=1,NRD
    DO 19 J=1,NAD
    DO 19 L=1,3
    DO 19M=1,3
    KT=3*(L-1)+M+9
    C(KT,I,J)=0.
C SET EXPECTED MAXIMUM RELATIVE VELOCITY IN COLLISIONS
    KV=3*(L-1)+M
    EM1=SPEC(L,1)
    EM2=SPEC(M,1)
    EMR=EM1*EM2/(EM1+EM2)
C
    IF(KS.EQ.1)GO TO 17
    TEM=FOE1(4,I,KR,KS)
    GO TO 18
17 TEM=FWP1(4,I,KR)
C

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18 RV=2./SQRT(PI*2.*BOLTZ*TEM/EMR) SIM03610
19 C(KV,I,J)=2.*RV SIM03620
C SIM03630
C MAXIMUM RELATIVE VELOCITY WILL BE RESET IF FASTER ENCOUNTERS OCCUR SIM03640
C SIM03650
C*****SIM03660
TIME=0. SIM03670
C LOOP OVER TIME INTERVALS.....BLOCK 10 SIM03680
DO 6000 JDTM = 1,NIS SIM03690
C SIM03700
C SIM03710
TIME=TIME+DTM SIM03720
C*****SIM03730
C MOVE ALL MOLECULES .....BLOCK 12 SIM03740
C MOVE MOLECULES OF SPECIES 1 SIM03750
C SIM03760
C SIM03770
DO 310 I1 = 1,NMOL1 SIM03780
C SKIP INACTIVE MOLECULES.....13 SIM03790
IF (P1(4,I1).EQ.-99.) GO TO 310 SIM03800
VX = P1(1,I1) SIM03810
VY = P1(2,I1) SIM03820
RX = P1(4,I1) SIM03830
T = P1(5,I1) SIM03840
TOLD=T SIM03850
C*****SIM03860
C FIND NEW COORDINATES.....BLOCK 14 SIM03870
XX=RX*COS(T)+VX*DTM SIM03880
YY=RX*SIN(T)+VY*DTM SIM03890
RNEW=SQRT(XX**2+YY**2) SIM03900
T=ATAN(YY/XX) SIM03910
C*****SIM03920
C FOR LAST SECTOR FIND COLLISIONS WITH THE WALL AND SAMPLE THEM SIM03930
C .....BLOCK 15,16 SIM03940
IF(ALFA.GT.DALFA)GO TO 301 SIM03950
IF(T.GT.0.)GO TO 301 SIM03960
DTR=DTM*T/(T-TOLD) SIM03970
C DTR IS THE TIME REMAINING AFTER A MOLECULE STRIKES THE WALL SIM03980
IF(DTR.LT.1E-10)DTR=1E-10 SIM03990
RW=RX+(RNEW-RX)*DTR/DTM SIM04000
IF(RW.LT.RMIN)RW=RMIN+DTR*.001 SIM04010
IF(RW.GT.RMAX)RW=RMAX-DTR*.001 SIM04020
LOC=RW/DR+1. SIM04030
C SIM04040
C COUNT COLLISIONS WITH THE WALL(MUST BE WAIGHTED =*DFI) SIM04050
SS(1,LOC,JSAMP)=SS(1,LOC,JSAMP)+1 SIM04060
C SIM04070
C SET VELOCITY AND LOCATION AFTER A MOLECULE STRIKES THE WALL SIM04080
302 CALL RANDU(P) SIM04090
B=VWM1*SQRT(-ALOG(P)) SIM04100
CALL RANDU(P) SIM04110
BB=2.*PI*P SIM04120
P1(1,NADR1)=B*COS(BB) SIM04130
VX=B*COS(BB) SIM04140
P1(2,NADR1)=B*SIN(BB) SIM04150
VY=B*SIN(BB) SIM04160
CALL RANDU(P) SIM04170
P1(3,NADR1)=VWM1*SQRT(-ALOG(P)) SIM04180
XX=RX*COS(T)+VX*DTR SIM04190
YY=RX*SIN(T)+VY*DTR SIM04200
RNEW=SQRT(XX**2+YY**2) SIM04210
T=ATAN(YY/XX) SIM04220
C SIM04230
C*****SIM04240
C DEACTIVATE ALL MOLECULES THAT MOVED OUT.....BLOCK 17 SIM04250
301 IF(RNEW.LT.RMAX.AND.RNEW.GT.RMIN.AND.T.LT.TMAX.AND.T.GT.TMIN)GO TO SIM04260
*303 SIM04270
RI=(RNEW-RMIN)/DR+.999999 SIM04280
IR=RI SIM04290
TI=(TMAX-T)/DDALFA+.999999 SIM04300
IT=TI SIM04310
C SIM04320

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IF(IT.LE.0)GO TO 304	SIM04330
IF(IT.GT.NAD)GO TO 305	SIM04340
IF(IR.GT.NRD)GO TO 306	SIM04350
C COUNT S DIRECTION, SAMPLE .....18	SIM04360
FSN1(1,IT,KR)=FSN1(1,IT,KR)+1.	SIM04370
GO TO 309	SIM04380
C COUNT W DIRECTION, SAMPLE.....18	SIM04390
304 IF(IR.LE.0)IR=1	SIM04400
IF(IR.GT.NRD)IR=NRD	SIM04410
FWN1(1,IR)=FWN1(1,IR)+1.	SIM04420
GO TO 309	SIM04430
C COUNT E DIRECTION, SAMPLE.....18	SIM04440
305 IF(IR.LE.0)IR=1	SIM04450
IF(IR.GT.NRD)IR=NRD	SIM04460
FEN1(1,IR)=FEN1(1,IR)+1.	SIM04470
GO TO 309	SIM04480
C COUNT N DIRECTION, SAMPLE.....18	SIM04490
306 FNN1(1,IT,KR)=FNN1(1,IT,KR)+1.	SIM04500
C SET NEW VALUES IN THE MOLECULE TABLE	SIM04510
C	SIM04520
309 RNEW=-99.	SIM04530
303 P1(4,I1)=RNEW	SIM04540
P1(5,I1)=T	SIM04550
310 CONTINUE	SIM04560
C	SIM04570
C.....	SIM04580
C REPEAT PROCEDURE FOR SPECIES 2 MOLECULES.....BLOCKS 12 TO18	SIM04590
DO 320 I2=1,NMOL2	SIM04600
C SKIP INACTIVE MOLECULES	SIM04610
IF(P2(4,I2).EQ.-99.)GO TO 320	SIM04620
VX = P2(1,I2)	SIM04630
VY = P2(2,I2)	SIM04640
RX = P2(4,I2)	SIM04650
T = P2(5,I2)	SIM04660
TOLD=T	SIM04670
C	SIM04680
XX=RX*COS(T)+VX*DTM	SIM04690
YY=RX*SIN(T)+VY*DTM	SIM04700
RNEW=SQRT(XX**2+YY**2)	SIM04710
T=ATAN(YY/XX)	SIM04720
C COLLISIONS WITH THE WALL	SIM04730
C.....	SIM04740
IF(ALFA.GT.DALFA)GO TO 311	SIM04750
IF(T.GT.0.)GO TO 311	SIM04760
DTR=DTM*T/(T-TOLD)	SIM04770
C DTR IS THE TIME REMAINING AFTER A MOLECULE STRIKES THE WALL	SIM04780
IF(DTR.LT.1E-10)DTR=1E-10	SIM04790
RW=RX+(RNEW-RX)*DTR/DTM	SIM04800
IF(RW.LT.RMIN)RW=RW+DTR*.001	SIM04810
IF(RW.GT.RMAX)RW=RW-DTR*.001	SIM04820
LOC=RW/DR+1	SIM04830
C COUNT COLLISIONS WITH THE WALL (MUST BE WEIGHTED =*DFI)	SIM04840
SS(2,LOC,JSAMP)=SS(2,LOC,JSAMP)+1	SIM04850
C	SIM04860
C FIND THE NEW COORDINATES OF THE MOLECULE	SIM04870
312 CALL RANDU(P)	SIM04880
B=VWM2*SQRT(-ALOG(P))	SIM04890
CALL RANDU(P)	SIM04900
BB=2.*PI*P	SIM04910
P2(1,NADR2)=B*COS(BB)	SIM04920
VX=B*COS(BB)	SIM04930
P2(2,NADR2)=B*SIN(BB)	SIM04940
VY=B*SIN(BB)	SIM04950
CALL RANDU(P)	SIM04960
XX=RX*COS(T)+VX*DTR	SIM04970
YY=RX*SIN(T)+VY*DTR	SIM04980
RNEW=SQRT(XX**2+YY**2)	SIM04990
T=ATAN(YY/XX)	SIM05000
C	SIM05010
C DEACTIVATE MOLECULES THAT MOVED OUT.COUNT FOR OUTPUT FLUX EVALUATION.	SIM05020
311 IF(RNEW.LT.RMAX.AND.RNEW.GT.RMIN.AND.T.LT.TMAX.AND.T.GT.TMIN)GO TO	SIM05030
*313	SIM05040



RI = (RNEW-RMIN)/DR+.999999	SIM05050
IR = RI	SIM05060
TI = (TMAX-T)/DDALFA+.999999	SIM05070
IT = TI	SIM05080
IF (IT.LE.0)GO TO 314	SIM05090
IF (IT.GT.NAD)GO TO 315	SIM05100
IF(IR.GT.NRD)GO TO 316	SIM05110
C COUNT S DIRECTION, SAMPLE.....18	SIM05120
FSN2(1,IT,KR)=FSN2(1,IT,KR)+1.	SIM05130
GO TO 319	SIM05140
C COUNT W DIRECTION, SAMPLE.....18	SIM05150
314 IF(IR.LE.0)IR=1	SIM05160
IF(IR.GT.NRD)IR=NRD	SIM05170
FWN2(1,IR)=FWN2(1,IR)+1.	SIM05180
GO TO 319	SIM05190
C COUNT E DIRECTION, SAMPLE.....18	SIM05200
315 IF(IR.LE.0)IR=1	SIM05210
IF(IR.GT.NRD)IR=NRD	SIM05220
FEN2(1,IR)=FEN2(1,IR)+1.	SIM05230
GO TO 319	SIM05240
C COUNT N DIRECTION, SAMPLE,.....18	SIM05250
316 FNN2(1,IT,KR)=FNN2(1,IT,KR)+1.	SIM05260
C	SIM05270
319 RNEW = -99.	SIM05280
C.....19	SIM05290
313 P2(4,I2)= RNEW	SIM05300
P2(5,I2)= T	SIM05310
320 CONTINUE	SIM05320
C	SIM05330
C*****	SIM05340
C NOW NEW MOLECULES WILL BE INTRODUCED	SIM05350
C TOTAL JET FLUX INTO THE REGION WAS DETERMINED BY F1(I) MOL./SEC.	SIM05360
C OR FWP1(I,KR),FWP2(I,KR) FOR EACH SPECIES	SIM05370
C NUMBER OF MOLECULES TO BE ACTIVATED PER DTM IS F(I)*DTM PER CELL.	SIM05380
C	SIM05390
C*****	SIM05400
DO 390 I=1,NRD	SIM05410
C INPUT 'W' MOLECULES	SIM05420
IF(KS.EQ.1)GO TO 322	SIM05430
ANEW1=FOE1(1,I,KR,KS-1)	SIM05440
ANEW2=FOE2(1,I,KR,KS-1)	SIM05450
GO TO 323	SIM05460
322 ANEW1=FWP1(1,I,KR)*DTM.	SIM05470
ANEW2=FWP2(1,I,KR)*DTM	SIM05480
323 NEW1=ANEW1	SIM05490
NEW2=ANEW2	SIM05500
REM1=ANEW1-NEW1	SIM05510
CALL RANDU(P)	SIM05520
IF(REM1.GT.P)NEW1=NEW1+1	SIM05530
REM2=ANEW2-NEW2	SIM05540
CALL RANDU(P)	SIM05550
IF(REM2.GT.P)NEW2=NEW2+1	SIM05560
C ACTIVATE NEW INPUT MOLECULES	SIM05570
C TIME, LOCATION AND VELOCITY COMP. OF NEW MOLs. ARE RANDOM FUNCTIONS	SIM05580
C SPECIES 1.....	SIM05590
IF(NEW1.LT.1)GO TO 341	SIM05600
C	SIM05610
DO 340 I1=1,NEW1	SIM05620
CALL RANDU(P)	SIM05630
ATIME=P*DTM	SIM05640
CALL RANDU(P)	SIM05650
RSTART=(FLOAT(I-1)+P)*DR+RMIN	SIM05660
VELX=FWP1(2,I,KR)	SIM05670
VELY=FWP1(3,I,KR)	SIM05680
TEM=FWP1(4,I,KR)	SIM05690
C THERMAL VELOCITY	SIM05700
VTER1=SQRT(2.*BOLTZ*TEM/SPEC(1,1))	SIM05710
C TEMPERATURE IS TRANSFERRED THROUGH FWP1(4,I,KR)	SIM05720
XX=RSTART*COS(ALFA)+VELX*ATIME	SIM05730
YY=RSTART*SIN(ALFA)+VELY*ATIME	SIM05740
RX=SQRT(XX**2+YY**2)	SIM05750
T=ATAN(YY/XX)	SIM05760

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      IF(RX.LT.RMIN.OR.RX.GT.RMAX)GO TO 340
      IF(T.LT.TMIN.OR.T.GT.TMAX)GO TO 340
C FOR MORE ACCURATE CALCULATION SET THESE MOLECULES IN OUTPUT FLOWS
C
C DEFINE THE VELOCITY COMPONENTS
      CALL RANDU(P)
      B=VTER1*SQRT(-ALOG(P))
      CALL RANDU(P)
      BB=2.*PI*P
      VELX=VELX+B*COS(BB)
      VELY=VELY+B*SIN(BB)
      CALL RANDU(P)
      VELZ=VTER1*SQRT(-ALOG(P))
C DEFINE THE NEW MOLECULE TO BE ACTIVATED
      DO 325 IACT=1,NMOL1
      IF(P1(4,IACT).EQ.-99.)GO TO 326
325 CONTINUE
C
C IF THERE IS NO ROOM FOR ADDITIONAL MOLECULE PRINT 'ALARM'
      IF(IACT.GE.NMOL1)GO TO 3004
326 P1(4,IACT)=RX
      P1(5,IACT)=T
      P1(1,IACT)=VELX
      P1(2,IACT)=VELY
      P1(3,IACT)=VELZ
340 CONTINUE
C.....
C REPEAT PROCEDURE FOR SPECIES 2
341 IF(NEW2.LT.1)GO TO 351
      DO 350 I2=1,NEW2
      CALL RANDU(P)
      ATIME=P*DTM
      CALL RANDU(P)
      RSTART=RMIN+(FLOAT(I-1)+P)*DR
      VELX=FWP2(2,I,KR)
      VELY=FWP2(3,I,KR)
      TEM=FWP2(4,I,KR)
C THERMAL VELOCITY
      VTER2=SQRT(2.*BOLTZ*TEM/SPEC(2,1))
      XX=RSTART*COS(ALFA)+VELX*ATIME
      YY=RSTART*SIN(ALFA)+VELY*ATIME
      RX=SQRT(XX**2+YY**2)
      T=ATAN(YY/XX)
      IF(RX.LT.RMIN.OR.RX.GT.RMAX)GO TO 350
      IF(T.LT.TMIN.OR.T.GT.TMAX)GO TO 350
C FOR MORE ACCURATE RESULTS SET THESE MOLECULES IN OUTPUT FLOWS
C
C DEFINE VELOCITY COMPONENTS
      CALL RANDU(P)
      B=VTER2*SQRT(-ALOG(P))
      CALL RANDU(P)
      BB=2.*PI*P
      VELX=VELX+B*COS(BB)
      VELY=VELY+B*SIN(BB)
      CALL RANDU(P)
      VELZ=VTER2*SQRT(-ALOG(P))
C FIND A NEW MOLECULE TO BE ACTIVATED
      DO 345 IACT=1,NMOL2
      IF (P2(4,IACT).EQ.-99.)GO TO 346
345 CONTINUE
C IF THERE IS NO PLACE THEN PRINT ALARM
      IF(IACT.EQ.NMOL2)GO TO 3004
346 P2(4,IACT)=RX
      P2(5,IACT)=T
      P2(1,IACT)=VELX
      P2(2,IACT)=VELY
      P2(3,IACT)=VELZ
350 CONTINUE
C
C REPEAT PROCEDURE FOR SPEC-3
C.....
C INPUT E MOLECULES

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SIM0577
SIM0578
SIM0579
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SIM0590
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SIM0646
SIM0647
SIM0648

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351	IF (TMIN.LE.0.)GO TO 390	SIM06490
C		SIM06500
C		SIM06510
	ANEW1=FOW1(1,I,KR,KS+1)	SIM06520
C		SIM06530
C		SIM06540
	NEW1=ANEW1	SIM06550
	REM=ANEW1-NEW1	SIM06560
	CALL RANDU(P)	SIM06570
	IF(P.LT.REM)NEW1=NEW1+1	SIM06580
C		SIM06590
	ANEW2=FOW2(1,I,KR,KS+1)	SIM06600
	NEW2=ANEW2	SIM06610
	REM=ANEW2-NEW2	SIM06620
	CALL RANDU(P)	SIM06630
	IF(P.LT.REM)NEW2=NEW2+1	SIM06640
C		SIM06650
C		SIM06660
	IF(NEW1.LE.0)GO TO 370	SIM06670
C		SIM06680
	DO 362 I1=1,NEW1	SIM06690
	CALL RANDU(P)	SIM06700
	T=TMIN+DDALFA*P	SIM06710
	CALL RANDU(P)	SIM06720
	RNEW=RMIN+(FLOAT(I-1)+P)*DR	SIM06730
	CALL RANDU(P)	SIM06740
	TEM=FOW1(4,I,KR,KS+1)	SIM06750
	VTER1=SQRT(2.*BOLTZ*TEM/SPEC(1,1))	SIM06760
	B=VTER1*SQRT(-ALOG(P))	SIM06770
	CALL RANDU(P)	SIM06780
	BB=2.*PI*P	SIM06790
	VELX=FOW1(2,I,KR,KS+1)+B*COS(BB)	SIM06800
	VELY=FOW1(3,I,KR,KS+1)+B*SIN(BB)	SIM06810
	CALL RANDU(P)	SIM06820
	VELZ=VTER1*SQRT(-ALOG(P))	SIM06830
C	FIND A NEW MOLECULE TO BE ACTIVATED	SIM06840
	DO 364 IACT=1,NMOL1	SIM06850
	IF(P1(4,IACT).EQ.-99.)GO TO 366	SIM06860
	364 CONTINUE	SIM06870
C	IF THERE IS NO ROOM FOR ADDITIONAL MOLECULE THEN PRINT 'ALARM'	SIM06880
	IF(IACT.EQ.NMOL1)GO TO 3004	SIM06890
366	P1(4,IACT)=RNEW	SIM06900
	P1(5,IACT)=T	SIM06910
	P1(1,IACT)=VELX	SIM06920
	P1(2,IACT)=VELY	SIM06930
362	P1(3,IACT)=VELZ	SIM06940
370	CONTINUE	SIM06950
C		SIM06960
C	REPEAT FOR SPECIES 2	SIM06970
	IF(NEW2.LE.0)GO TO 390	SIM06980
	DO 382 I2=1,NEW2	SIM06990
	CALL RANDU(P)	SIM07000
	T=TMIN+DDALFA*P	SIM07010
	CALL RANDU(P)	SIM07020
	RNEW=RMIN+(FLOAT(I-1)+P)*DR	SIM07030
	CALL RANDU(P)	SIM07040
	TEM=FOW2(4,I,KR,KS+1)	SIM07050
	VTER2=SQRT(2.*BOLTZ*TEM/SPEC(2,1))	SIM07060
	B=VTER2*SQRT(-ALOG(P))	SIM07070
	CALL RANDU(P)	SIM07080
	BB=2.*PI*P	SIM07090
	VELX=FOW2(2,I,KR,KS+1)+B*COS(BB)	SIM07100
	VELY=FOW2(3,I,KR,KS+1)+B*COS(BB)	SIM07110
	CALL RANDU(P)	SIM07120
	VELZ=VTER2*SQRT(-ALOG(P))	SIM07130
C	FIND A NEW MOLECULE TO BE ACTIVATED	SIM07140
	DO 384 IACT=1,NMOL2	SIM07150
	IF(P2(4,IACT).EQ.-99)GO TO 386	SIM07160
	384 CONTINUE	SIM07170
C	IF THERE IS NO ROOM FOR ADDITIONAL MOLECULE THEN PRINT ALARM	SIM07180
	IF(IACT.EQ.NMOL2) GO TO 3004	SIM07190
386	P2(4,IACT)=RNEW	SIM07200



P2(5, IACT)=T	SIM07210
P2(1, IACT)=VELX	SIM07220
P2(2, IACT)=VELY	SIM07230
382 P2(3, IACT)=VELZ	SIM07240
C.....	SIM07250
C REPEAT FOR SPEC-3	SIM07260
C	SIM07270
390 CONTINUE	SIM07280
C.....	SIM07290
C INPUT 'S' MOLECULES	SIM07300
DO 460 J=1, NAD	SIM07310
IF(KR.LT.2)GO TO 460	SIM07320
ANEW1=FNN1(1, J, KR-1)	SIM07330
IF(ANEW1.LE..00001)GO TO 420	SIM07340
NEW1=ANEW1	SIM07350
REM=ANEW1-NEW1	SIM07360
CALL RANDU(P)	SIM07370
IF(P.LT.REM)NEW1=NEW1+1	SIM07380
DO 402 I1=1, NEW1	SIM07390
CALL RANDU(P)	SIM07400
T=TMIN+DDALFA*(P+FLOAT(J-1))	SIM07410
CALL RANDU(P)	SIM07420
RNEW=RMIN+P*DR	SIM07430
CALL RANDU(P)	SIM07440
TEM=FNN1(4, J, KR-1)	SIM07450
VTER1=SQRT(2.*BOLTZ*TEM/SPEC(1,1))	SIM07460
B=VTER1*SQRT(-ALOG(P))	SIM07470
CALL RANDU(P)	SIM07480
BB=2.*PI*P	SIM07490
VELX=FNN1(2, J, KR-1)+B*COS(BB)	SIM07500
VELY=FNN1(3, J, KR-1)+B*SIN(BB)	SIM07510
CALL RANDU(P)	SIM07520
VELZ=VTER1*SQRT(-ALOG(P))	SIM07530
C FIND A NEW MOLECULE TO BE ACTIVATED	SIM07540
DO 404 IACT=1, NMOL1	SIM07550
IF(P1(4, IACT).EQ.-99.)GO TO 406	SIM07560
404 CONTINUE	SIM07570
IF(IACT.EQ.NMOL1)GO TO 3004	SIM07580
406 P1(4, IACT)=RNEW	SIM07590
P1(5, IACT)=T	SIM07600
P1(1, IACT)=VELX	SIM07610
P1(2, IACT)=VELY	SIM07620
402 P1(3, IACT)=VELZ	SIM07630
420 CONTINUE	SIM07640
C REPEAT FOR SPECIES 2	SIM07650
ANEW2=FNN2(1, J, KR-1)	SIM07660
IF(ANEW2.LT..00001)GO TO 440	SIM07670
NEW2=ANEW2	SIM07680
REM=ANEW2-NEW2	SIM07690
CALL RANDU(P)	SIM07700
IF(P.LT.REM)NEW2=NEW2+1	SIM07710
DO 422 I2=1, NEW2	SIM07720
CALL RANDU(P)	SIM07730
T=TMIN+DDALFA*(P+FLOAT(J-1))	SIM07740
CALL RANDU(P)	SIM07750
RNEW=RMIN+P*DR	SIM07760
CALL RANDU(P)	SIM07770
TEM=FNN2(4, J, KR-1)	SIM07780
VTER2=SQRT(2.*BOLTZ*TEM/SPEC(2,1))	SIM07790
B=VTER2*SQRT(-ALOG(P))	SIM07800
CALL RANDU(P)	SIM07810
BB=2.*PI*P	SIM07820
VELX=FNN2(2, J, KR-1)+B*COS(BB)	SIM07830
VELY=FNN2(3, J, KR-1)+B*SIN(BB)	SIM07840
CALL RANDU(P)	SIM07850
VELZ=VTER2*SQRT(-ALOG(P))	SIM07860
C FIND A NEW MOLECULE TO BE ACTIVATED	SIM07870
DO 424 IACT=1, NMOL2	SIM07880
IF(P2(4, IACT).EQ.-99)GO TO 426	SIM07890
424 CONTINUE	SIM07900
IF(IACT.EQ.NMOL2)GO TO 3004	SIM07910
426 P2(4, IACT)=RNEW	SIM07920

P2(5, IACT)=T	SIM07930
P2(1, IACT)=VELX	SIM07940
P2(2, IACT)=VELY	SIM07950
422 P2(3, IACT)=VELZ	SIM07960
440 CONTINUE	SIM07970
C.....	SIM07980
C REPEAT FOR SPEC-3	SIM07990
C.....	SIM08000
C INPUT 'N' MOLECULES.....	SIM08010
C SPEC-1	SIM08020
ANEW1=FSN1(1, J, KR+1)	SIM08030
IF(ANEW1.LE..00001)GO TO 450	SIM08040
NEW1=ANEW1	SIM08050
REM=ANEW1-NEW1	SIM08060
CALL RANDU(P)	SIM08070
IF (P.LT.REM)NEW1=NEW1+1	SIM08080
DO 442 I1=1, NEW1	SIM08090
CALL RANDU(P)	SIM08100
T=TMIN +DDALFA*(P+FLOAT(J-1))	SIM08110
CALL RANDU(P)	SIM08120
RNEW=RMAX-P*DR	SIM08130
CALL RANDU(P)	SIM08140
TEM=FSN1(4, J, KR+1)	SIM08150
VTER1=SQRT(2.*BOLTZ*TEM/SPEC(1,1))	SIM08160
B=VTER1*SQRT(-ALOG(P))	SIM08170
CALL RANDU(P)	SIM08180
BB=2.*PI*P	SIM08190
VELX=FSN1(2, J, KR+1)+B*COS(BB)	SIM08200
VELY=FSN1(3, J, KR+1)+B*SIN(BB)	SIM08210
CALL RANDU(P)	SIM08220
VELZ=VTER1*SQRT(-ALOG(P))	SIM08230
C FIND A NEW MOLECULE TO BE ACTIVATED	SIM08240
DO 444 IACT=1, NMOL1	SIM08250
IF(P1(4, IACT).EQ.-99.)GO TO 446	SIM08260
444 CONTINUE	SIM08270
IF(IACT.EQ.NMOL1)GO TO 3004	SIM08280
446 P1(4, IACT)=RNEW	SIM08290
P1(5, IACT)=T	SIM08300
P1(1, IACT)=VELX	SIM08310
P1(2, IACT)=VELY	SIM08320
442 P1(3, IACT)=VELZ	SIM08330
450 CONTINUE	SIM08340
C.....	SIM08350
C INPUT N MOLECULES SPEC-2	SIM08360
ANEW2=FSN2(1, J, KR+1)	SIM08370
IF(ANEW2.LE..00001)GO TO 460	SIM08380
NEW2=ANEW2	SIM08390
REM=ANEW2-NEW2	SIM08400
CALL RANDU(P)	SIM08410
IF(P.LT.REM)NEW2=NEW2+1	SIM08420
DO 452 I2=1, NEW2	SIM08430
CALL RANDU(P)	SIM08440
T=TMIN+DDALFA*(P+FLOAT(J-1))	SIM08450
CALL RANDU(P)	SIM08460
RNEW=RMAX-P*DR	SIM08470
CALL RANDU(P)	SIM08480
TEM=FSN2(4, J, KR+1)	SIM08490
VTER2=SQRT(2.*BOLTZ*TEM/SPEC(2,1))	SIM08500
B=VTER2*SQRT(-ALOG(P))	SIM08510
CALL RANDU(P)	SIM08520
BB=2*PI*P	SIM08530
VELX=FSN2(2, J, KR+1)+B*COS(BB)	SIM08540
VELY=FSN2(3, J, KR+1)+B*SIN(BB)	SIM08550
CALL RANDU(P)	SIM08560
VELZ=VTER2*SQRT(-ALOG(P))	SIM08570
C FIND A NEW MOLECULE TO BE ACTIVATED	SIM08580
DO 454 IACT=1, NMOL2	SIM08590
IF(P2(4, IACT).EQ.-99)GO TO 456	SIM08600
454 CONTINUE	SIM08610
IF(IACT.EQ.NMOL2)GO TO 3004	SIM08620
456 P2(4, IACT)=RNEW	SIM08630
P2(5, IACT)=T	SIM08640

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      P2(1,IACT)=VELX
      P2(2,IACT)=VELY
452  P2(3,IACT)=VELZ
460  CONTINUE
C
C
C*****
C*****
C REARRANGE MOLECULES IN THEIR CELLS
C INITIALIZATION
  5000 CONTINUE
C*****
      KIP=NMOL1+NMOL2+NMOL3
      DO 1001 KADD=1,KIP
1001  IP(KADD)=0
      KADD=0
      N1A=0
      N2A=0
      N3A=0
C N1A,N2A,N3A ARE THE NUMBER OF ACTIVE MOLECULES (COUNTED NEXT)
C
      DO 1100 I=1,NRD
      DO 1100 J=1,NAD
C
      DO 1005 K=1,4
1005  IC(K,I,J)=0
C
C SET SPECIES 1
      N1C=0
      N2C=0
      N3C=0
      NTC=0
      DO 1020 K1=1,NMOL1
      IF(P1(4,K1).EQ.-99.)GO TO 1020
      RLOC=ABS(C(19,I,J)-P1(4,K1))
      TLOC=ABS(C(20,I,J)-P1(5,K1))
      IF(RLOC.GT.DR*.5)GO TO 1020
      IF(TLOC.GT.DDALFA*.5)GO TO 1020
C
      N1C=N1C+1
      NTC=NTC+1
      N1A=N1A+1
      IF(NTC.EQ.1)IC(4,I,J)=KADD
      IC(1,I,J)=IC(1,I,J)+1
      KADD=KADD+1
      IP(KADD)=K1
1020  CONTINUE
C SET SPEC=2 MOLECULES.....
      DO 1040 K2=1,NMOL2
      IF(P2(4,K2).EQ.-99.)GO TO 1040
      RLOC=ABS(C(19,I,J)-P2(4,K2))
      TLOC=ABS(C(20,I,J)-P2(5,K2))
      IF(RLOC.GT.DR*.5)GO TO 1040
      IF(TLOC.GT.DDALFA*.5)GO TO 1040
C
      N2A=N2A+1
      NTC=NTC+1
      IF(NTC.EQ.1)IC(4,I,J)=KADD
      IC(2,I,J)=IC(2,I,J)+1
      KADD=KADD+1
      IP(KADD)=K2
1040  CONTINUE
1100  CONTINUE
      WRITE(6,1021)N1A,N2A
1021  FORMAT('  NUMBER OF ACTIVE MOLECULES SPEC1= ',I5,' SPEC2=',I5/)
C
C
C*****
C CALCULATE COLLISIONS.....BLOCK
C EXTERNAL LOOPS ARE OVER CELLS I,J
C*****
C

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DO 999 I=1,NRD	SIM09370
DO 999 J=1,NAD	SIM09380
999 NCOL(I,J)=0	SIM09390
C	SIM09400
DO 900 I=1,NRD	SIM09410
DO 900 J=1,NAD	SIM09420
C	SIM09430
NM(1)=IC(1,I,J)	SIM09440
NM(2)=IC(2,I,J)	SIM09450
NM(3)=IC(3,I,J)	SIM09460
NTOT=NM(1)+NM(2)+NM(3)	SIM09470
DO 900 L=1,3	SIM09480
DO 900 M=1,3	SIM09490
C NUMBER OF SPECIES IN THE PROGRAM IS 2 (3)	SIM09500
KV=(L-1)*3+M	SIM09510
KT=KV+9	SIM09520
C KV,KT ARE THE ADDRESSES OF RELATIVE VELOCITY AND COLLISION TIMERS	SIM09530
920 IF(C(KT,I,J).GE.TIME) GO TO 900	SIM09540
C C(KT,I,J) IS THE INTEGRATED TIME FOR L-M COLLISION	SIM09550
KSEL=0	SIM09560
KREJ=0	SIM09570
IF(NM(L).GT.1.AND.NM(M).GT.1)GO TO 912	SIM09580
C NO COLLISIONS ARE CALCULATED IF THERE ARE NO MOLECULES	SIM09590
911 C(KT,I,J)=C(KT,I,J)+DTM	SIM09600
GO TO 900	SIM09610
C SELECT NOW THE MOLECULES FOR COLLISION	SIM09620
912 IF (KSEL.GE.100)GO TO 911	SIM09630
CALL RANDU(P)	SIM09640
MOL1=P*NM(L)+.999999	SIM09650
IF(MOL1.EQ.0)MOL1=1	SIM09660
C	SIM09670
CALL RANDU(P)	SIM09680
MOL2=P*NM(M)+.999999	SIM09690
IF(MOL2.EQ.0)MOL2=1	SIM09700
C	SIM09710
KSEL=KSEL+1	SIM09720
C	SIM09730
C CHECK IF THE SAME MOLECULE HAS BEEN SELECTED TWICE	SIM09740
IF(L.EQ.M.AND.MOL1.EQ.MOL2)GO TO 912	SIM09750
C.....	SIM09760
C FIND THE ACTUAL ADDRESSES OF THE SELECTED MOLECULES	SIM09770
IF(L.EQ.1)K1=0	SIM09780
IF(L.EQ.2)K1=NM(1)	SIM09790
IF(L.EQ.3)K1=NM(1)+NM(2)	SIM09800
IF(M.EQ.1)K2=0	SIM09810
IF(M.EQ.2)K2=NM(1)	SIM09820
IF(M.EQ.3)K2=NM(1)+NM(2)	SIM09830
KAD1=MOL1+K1+IC(4,I,J)	SIM09840
KAD2=MOL2+K2+IC(4,I,J)	SIM09850
C KAD1,KAD2 ARE THE LOCATION OF SELECTED MOLECULES IN IP( )	SIM09860
MAD1=IP(KAD1)	SIM09870
MAD2=IP(KAD2)	SIM09880
C MAD1,MAD2 ARE THE ACTUAL ADDRESSES OF THE SELECTED MOLECULES (THE	SIM09890
C INDICATION OF WHAT SPECIES THEY ARE HAS BEEN DEFINED HERE)	SIM09900
DO 930 N=1,3	SIM09910
IF(L.EQ.1)VN1=P1(N,MAD1)	SIM09920
IF(L.EQ.2)VN1=P2(N,MAD1)	SIM09930
IF(L.EQ.3)VN1=P3(N,MAD1)	SIM09940
C	SIM09950
IF(M.EQ.1)VN2=P1(N,MAD2)	SIM09960
IF(M.EQ.2)VN2=P2(N,MAD2)	SIM09970
IF(M.EQ.3)VN2=P3(N,MAD2)	SIM09980
C	SIM09990
930 VRC(N)=VN1-VN2	SIM10000
C VRC(3) CONTAIN THE THREE RELATIVE VELOCITY COMPONENTS	SIM10010
VR=SQRT(VRC(1)*VRC(1)+VRC(2)*VRC(2)+VRC(3)*VRC(3))	SIM10020
C VR IS THE RELATIVE SPEED IN A SPECIFIC COLLISION	SIM10030
IF(C(KV,I,J).LT.VR)C(KV,I,J)=VR	SIM10040
C LAST STATEMENT RESETS THE MAXIMUM RELATIVE VELOCITY FOR FURTHER	SIM10050
C CALCULATIONS	SIM10060
C	SIM10070
IF(KREJ.GT.100)GO TO 911	SIM10080

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C      CALL RANDU(P)
      AVR=VR/C(KV,I,J)
      KREJ=KREJ+1
      IF(AVR.LT.P)GO TO 912
C LAST STATEMENT REJECTS THE CALCULATED COLLISION
C
C NOW A SPECIES L-M COLLISION HAS BEEN SELECTED
C
C CALCULATE NOW THE PROBABILITY THAT SUCH A COLLISION WILL BE COUNTED
C FOR THE L AND M SPECIES RESPECTIVELY
      LP=1
      LM=1
      CALL RANDU(P)
      ANM=FLOAT(NM(L))/FLOAT(NM(M))
      IF(ANM.GT.1.)GO TO 950
      IF(ANM.LT.P)MP=0
      GO TO 955

C
C
950  ANM=1./ANM
      IF(ANM.LT.P)LP=0

C
C
955  CXS=PI*(SPEC(L,2)+SPEC(M,2))*2/4.
      ALP=LP
      ALM=LM

C      VOLUME=REG(4,KR,KS)*VOL(I)
C.....CHECK
      DNL=FLOAT(NM(L))/VOLUME
      DNM=FLOAT(NM(M))/VOLUME
C USE EQ.10.3
      TOC(L,M)=ALP/(CXS*DNM*VR*NM(L))+ALM/(CXS*DNL*VR*NM(M))
C
C SET THIS VALUE INTO C(KT,I,J)
      C(KT,I,J)=C(KT,I,J)+TOC(L,M)
C.....
C SAMPLE THIS COLLISION
      NCOL(I,J)=NCOL(I,J)+1
C
C FIND RELATIVE MASSES
      CALL RANDU(P)
      BB=1.-2.*P
      AA=SQRT(1.-BB*BB)
      VRC(1)=BB*VR
      CALL RANDU(P)
      BB=2.*PI*P
      VRC(2)=AA*COS(BB)*VR
      VRC(3)=AA*SIN(BB)*VR
C
C FIND RELATIVE MASSES
      SM=SPEC(L,1)+SPEC(M,1)
      RML=SPEC(L,1)/SM
      RMM=SPEC(M,1)/SM
C
C*****
C CALCULATE HERE THE ACTUAL ADDRESSES OF COLLIDING MOLECULES AND SET
C THEIR NEW VELOCITIES (NEW VELOCITY COMPONENTS ARE ADDED TO THE
C VELOCITY OF THE CENTER OF MASS VCCM)
      DO 960 N=1,3
        IF(L.EQ.1)V1=P1(N,MAD1)
        IF(L.EQ.2)V1=P2(N,MAD1)
C      IF(L.EQ.3)V1=P3(N,MAD1)
        IF(M.EQ.1)V2=P1(N,MAD2)
        IF(M.EQ.2)V2=P2(N,MAD2)
C      IF(M.EQ.3)V2=P3(N,MAD2)
C
C      VCCM=RML*V1+RMM*V2
      VCCM1=VCCM+VRC(N)*RMM
      VCCM2=VCCM-VRC(N)*RML
C

```

```

SIM10091
SIM10101
SIM10111
SIM10121
SIM10131
SIM10141
SIM10151
SIM10161
SIM10171
SIM10181
SIM10191
SIM10201
SIM10211
SIM10221
SIM10231
SIM10241
SIM10251
SIM10261
SIM10271
SIM10281
SIM10291
SIM10301
SIM10311
SIM10321
SIM10331
SIM10341
SIM10351
SIM10361
SIM10371
SIM10381
SIM10391
SIM10401
SIM10411
SIM10421
SIM10431
SIM10441
SIM10451
SIM10461
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SIM10481
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SIM10521
SIM10531
SIM10541
SIM10551
SIM10561
SIM10571
SIM10581
SIM10591
SIM10601
SIM10611
SIM10621
SIM10631
SIM10641
SIM10651
SIM10661
SIM10671
SIM10681
SIM10691
SIM10701
SIM10711
SIM10721
SIM10731
SIM10741
SIM10751
SIM10761
SIM10771
SIM10781
SIM10791
SIM10801

```

C	CHANGE IN VELOCITY IS INPUTED ONLY IF PROBABILITY OF COLLISION.GT.1	SIM10810
	IF(LP.NE.1)GO TO 961	SIM10820
	IF(L.EQ.1)P1(N,MAD1)=VCCM1	SIM10830
	IF(L.EQ.2)P2(N,MAD1)=VCCM1	SIM10840
C	IF(L.EQ.3)P3(N,MAD1)=VCCM1	SIM10850
C		SIM10860
	961 IF(LM.NE.1)GO TO 960	SIM10870
	IF(M.EQ.1)P1(N,MAD2)=VCCM2	SIM10880
	IF(M.EQ.2)P2(N,MAD2)=VCCM2	SIM10890
C	IF(M.EQ.3)P3(N,MAD2)=VCCM2	SIM10900
C		SIM10910
	960 CONTINUE	SIM10920
	GO TO 920	SIM10930
	900 CONTINUE	SIM10940
	WRITE(6,4545)TOC(1,1),TOC(1,2),TOC(2,1),TOC(2,2),TIME	SIM10950
	4545 FORMAT(' TIME',5E10.3)	SIM10960
C		SIM10970
	C*****END OF COLLISIONS*****	SIM10980
	C NOW NEW TEMPERATURES MAY BE CALCULATED IN EACH CELL	SIM10990
	C AVERAGE VELOCITY IN CELLS	SIM11000
C		SIM11010
	WRITE(6,4547)	SIM11020
	4547 FORMAT('0 I J N1E N2E TEMPR1 TEMPR2 VAVX1	SIM11030
	1 VAVY1 VAVX2 VAVY2 NCOL',/)	SIM11040
C		SIM11050
	DO 1110 I=1,NRD	SIM11060
	DO 1110 J=1,NAD	SIM11070
	N1E=IC(1,I,J)	SIM11080
	N2E=IC(2,I,J)	SIM11090
	KAD1=IC(4,I,J)	SIM11100
	KAD2=IC(4,I,J)+N1E	SIM11110
	IF(N1E.LT.1)GO TO 1112	SIM11120
C		SIM11130
	VX1=0.	SIM11140
	VX2=0.	SIM11150
	VY1=0.	SIM11160
	VY2=0.	SIM11170
C		SIM11180
	DO 1111 IM=1,N1E	SIM11190
	MAD1=KAD1+IM	SIM11200
	IAD=IP(MAD1)	SIM11210
	VX1=VX1+P1(1,IAD)	SIM11220
	1111 VY1=VY1+P1(2,IAD)	SIM11230
	1112 IF(N2E.LE.0)GO TO 1110	SIM11240
C		SIM11250
	DO 1115 IM=1,N2E	SIM11260
	MAD2=MAD1+IM	SIM11270
	IAD=IP(MAD2)	SIM11280
	VX2=VX2+P2(1,IAD)	SIM11290
	1115 VY2=VY2+P2(2,IAD)	SIM11300
C		SIM11310
	C THE AVERAGE VELOCITIES IN THE CELL WILL RESULT-	SIM11320
	VAVX1=VX1/N1E	SIM11330
	VAVY1=VY1/N1E	SIM11340
	VAVX2=VX2/N2E	SIM11350
	VAVY2=VY2/N2E	SIM11360
	IF(I.NE.1)GO TO 1116	SIM11370
C		SIM11380
	FSN1(2,J,KR)=VAVX1+FSN1(2,J,KR)	SIM11390
	FSN1(3,J,KR)=VAVY1+FSN1(3,J,KR)	SIM11400
	FSN2(2,J,KR)=VAVX2+FSN2(2,J,KR)	SIM11410
	FSN2(3,J,KR)=VAVY2+FSN2(3,J,KR)	SIM11420
C		SIM11430
	1116 IF(I.NE.NRD)GO TO 1117	SIM11440
	FNN1(2,J,KR)=VAVX1+FNN1(2,J,KR)	SIM11450
	FNN1(3,J,KR)=VAVY1+FNN1(3,J,KR)	SIM11460
	FNN2(2,J,KR)=VAVX2+FNN2(2,J,KR)	SIM11470
	FNN2(3,J,KR)=VAVY2+FNN2(3,J,KR)	SIM11480
C		SIM11490
	1117 IF(J.NE.1)GO TO 1118	SIM11500
	FEN1(2,I)=VAVX1+FEN1(2,I)	SIM11510
	FEN1(3,I)=VAVY1+FEN1(3,I)	SIM11520



```

FEN2(2,I)=VAVX2+FEN2(2,I)
FEN2(3,I)=VAVY2+FEN2(3,I)
C
1118 IF(J.NE.NAD)GO TO 1119
      FWN1(2,I)=VAVX1+FWN1(2,I)
      FWN1(3,I)=VAVY1+FWN1(3,I)
      FWN2(2,I)=VAVX2+FWN2(2,I)
      FWN2(3,I)=VAVY2+FWN2(3,I)
1119 CONTINUE
C
C THERMAL VELOCITIES AND TEMPERATURES
      ENRG1=0.
      ENRG2=0.
      IF(N1E.LT.1)GO TO 1131
      DO 1130 IM=1,N1E
      MAD1=KAD1+IM
      IAD=IP(MAD1)
      CX1=P1(1,IAD)-VAVX1
      CY1=P1(2,IAD)-VAVY1
      CZ1=P1(3,IAD)
1130 ENRG1=ENRG1+CX1*CX1+CY1*CY1+CZ1*CZ1
1131 IF(N2E.LT.1)GO TO 1150
      DO 1140 IM=1,N2E
      MAD2=KAD2+IM
      IAD=IP(MAD2)
      CX2=P2(1,IAD)-VAVX2
      CY2=P2(2,IAD)-VAVY2
      CZ2=P2(3,IAD)
C
C
1140 ENRG2=ENRG2+CX2*CX2+CY2*CY2+CZ2*CZ2
C
1150 TEMPR1=(ENRG1/N1E)*SPEC(1,1)/(3.*BOLTZ)
      TEMPR2=(ENRG2/N2E)*SPEC(2,1)/(3.*BOLTZ)
C
      IF(I.NE.1)GO TO 1151
      FSN1(4,J,KR)=TEMPR1+FSN1(4,J,KR)
      FSN2(4,J,KR)=TEMPR2+FSN2(4,J,KR)
C
1151 IF(I.NE.NRD)GO TO 1152
      FNN1(4,J,KR)=TEMPR1+FNN1(4,J,KR)
      FNN2(4,J,KR)=TEMPR2+FNN2(4,J,KR)
C
1152 IF(J.NE.1)GO TO 1153
      FWN1(4,I)=TEMPR1+FWN1(4,I)
      FWN2(4,I)=TEMPR2+FWN2(4,I)
C
1153 IF(J.NE.NAD)GO TO 1154
      FEN1(4,I)=TEMPR1+FEN1(4,I)
      FEN2(4,I)=TEMPR2+FEN2(4,I)
1154 CONTINUE
C
      WRITE(6,4548)I,J,N1E,N2E,TEMPR1,TEMPR2,VAVX1,VAVY1,VAVX2,VAVY2,NCOSIM12050
      1L(I,J)
4548 FORMAT(' ',4I5,6F12.3,I7)
C
1110 CONTINUE
C
6000 CONTINUE
C CALCULATE AVERAGED PARAMETERS, WEIGHTED BY DFI
      DO 6010 I=1,NRD
      DO 6010 KPAR=1,4
      FOW1(KPAR,I,KR,KS)=FWN1(KPAR,I)/(NIS)
      FOW2(KPAR,I,KR,KS)=FWN2(KPAR,I)/(NIS)
      FOE1(KPAR,I,KR,KS)=FEN1(KPAR,I)/(NIS)
      FOE2(KPAR,I,KR,KS)=FEN2(KPAR,I)/(NIS)
6010
C
      DO 6020 J=1,NAD
      DO 6020 KPAR=1,4
      FSN1(KPAR,J,KR)=FSN1(KPAR,J,KR)/NIS
      FSN2(KPAR,J,KR)=FSN2(KPAR,J,KR)/NIS
      FNN1(KPAR,J,KR)=FNN1(KPAR,J,KR)/NIS

```

```

6020 FNN2(KPAR,J,KR)=FNN2(KPAR,J,KR)/NIS
C
C TO PREPARE FLOWS FOR THE NEXT REGION OR SECTOR, DIVIDE F(N) BY LOCAL DFI AND MULTIPLY BY NEXT DFI WHEN STARTING NEW REGION
C
C CALCULATE HERE THE MEAN FREE PATH AND STORE INTO REG(,,)
C
C STOP PROGRAM IF FLOW BECOMES COLLISIONLESS OR NUMBER DENSITY IS EQUAL TO THE AMBIENT NUMBER DENSITY. STORE FOE1,FOE2 IN A SEPARATE FILE TO BE USED IN A DIFFERENT PROGRAM.
C
C PRINT AVERAGED RESULTS
  WRITE(6,6029)
6029 FORMAT(' ',,////)
  WRITE(6,6030)NIS,KR
6030 FORMAT(' AVERAGED OUTPUT FLOWS AFTER',I5,' TIME INCREMENTS IN REGION',I5)
  WRITE(6,6031)
6031 FORMAT('0      I      FEN1(1,I)      FEN2(1,I)      FWN1(1,I)      FWN2(1,I)'
1)
  DO 6033 I=1,NRD
  WRITE(6,6032)I,FEN1(1,I),FEN2(1,I),FWN1(1,I),FWN2(1,I)
6032 FORMAT(' ',I5,4F13.3)
6033 CONTINUE
  WRITE(6,6034)
6034 FORMAT('0      J      FNN1(1,J,KR)      FNN2(1,J,KR)      FSN1(1,J,KR)      FSN2(1,J,KR)'
12(1,J,KR)')
  DO 6036 J=1,NAD
  WRITE(6,6037)J,FNN1(1,J,KR),FNN2(1,J,KR),FSN1(1,J,KR),FSN2(1,J,KR)
6037 FORMAT(' ',I5,4F15.5)
6036 CONTINUE
C
C*****
C
C START A NEW REGION
C   IF(KR.EQ.10) GO TO 7000
C   KR=KR+1
C   GO TO 2000
C
C*****
C
7000 CONTINUE
C
C*****
C
C FIND IF FLOW BECAME COLLISIONLESS IN THE WHOLE SECTOR
C IF POSITIVE, STORE FOE1,FOE2 IN A SEPARATE FILE AND STOP PROGRAM
C
C PREPARE DATA FOR THE NEXT SECTOR
C   KR=1
C   STOP PROGRAM IF KS WAS BOUNDED BY THE WALL
C   KS=KS+1
C   GO TO 1000
C
C*****
C
C IF THERE IS BACK FLOW (FWN1,FWN2) NONZERO CALCULATE NEXT ITERATION
C   ITER=ITER+1
C   KS=KR=1
C   GO TO 3000
C
C*****
C
  WRITE(6,6099)
6099 FORMAT('1 DATA FOR TEN MOLECULES SPEC.2')
  DO 3003 I=1,10
  WRITE(6,3002) P2(1,I),P2(2,I),P2(3,I),P2(4,I),P2(5,I)
3002 FORMAT(' ',4F10.3,E18.10)
3003 CONTINUE
  GO TO 3009
3004 WRITE(6,3005)

```



```
3005 FORMAT(' NO PLACE FOR ADDITIONAL MOLECULES')
3009 CONTINUE
      STOP
      END
```

C  
C

```
      SUBROUTINE RANDU(P)
      COMMON IX
      IY = IX*65539
      IF (IY) 5,6,6
5     IY = IY+2147483647+1
6     P = IY
      P = P*.4656613E-9
      IX = IY
      RETURN
      END
```

SIM12970  
SIM12980  
SIM12990  
SIM13000  
SIM13010  
SIM13020  
SIM13030  
SIM13040  
SIM13050  
SIM13060  
SIM13070  
SIM13080  
SIM13090  
SIM13100  
SIM13110  
SIM13120

## B.5 Program SIMUL - User's Guide

Preparation: Run AXSYM program. From its output data evaluate:

ALFA - The averaged angle for the continuum breakdown

( $P \sim 0.05$ )

TETA - Flow direction along the breakdown limit.

Flow parameters along this line - Pressure, temperature, velocity, mean free path.

Input Data:

Radius of nozzle ring	R1
Radial size of a cell	DR
Maximum radius in simulation	RP
Angle of breakdown limit	ALFA
Flow direction along (ALFA)	TETA
Averaged flow velocity along (ALFA)	Vo
Molecular weight of each species	Spec(I,1)
Molecular diameter of each species	Spec (I,2)
Mean free path along (ALFA)	FPM
Averaged Temperature (ALFA)	TEMP
Time increment	DTM
Number of time increments	NIS

Options

### a. Geometry

The program is designed to run for axisymmetric ring flow.

For a two dimensional planar flow the molecular motion, collisions and flow calculation remain unchanged. The flow cross section remain

unchanged. Instead of the angle DFI the two dimensional flow requires the definition of the width of each region (or cell). To keep the number of molecules within reasonable computational limits this size has to decrease in the same manner as DFI.

The part of the program which has to be changed for this purpose is lines 230 to 250.

b. Wall flux calculations:

The program may run for the whole molecular region resulting the flux towards the wall (this part needs additional debugging). However in order to make it more efficient we may stop the program at a sector where the flow becomes collisionless. The remaining part of the flow may be regarded either as collisionless or if we define a much larger size of cells we may calculate the molecular collisions on this basis.

This part need additional analysis and programming.

Execution Commands

Without additional changes the program runs under WATFIV compiler using the following command:

```
WATFI AXSYM * (XTYPE)
```

Further developments will be required to run the program for each sector separately and the output intermediate results on the mass storage.

## B.6 The Influence of the Ambient Gas

The temperature, pressure and density of the ambient gas are shown in Table 1. Because its temperature is much higher than in the jet gas, the thermal velocity of ambient gas molecules is much higher than jet molecules. the following contribution may be expected due to the ambient gas:

- a. Collisions between the "hot" ambient molecules with the "cold" jet molecules may cause an increase in the dissipation in the outer layer of the jet and increase in the flux towards the walls.
- b. For higher ambient pressures, those collisions become rare because of the low number density therefore, the influence of the collisions may decrease.

The only way to evaluate the influence of these two controversial factors is by an additional simulation program to be designed for this region. The following are the main factors to be included in this program:

### Boundary Conditions:

- a. The jet side: FOE1 and FOE2 obtained from the last simulated sector (SIMUL) supply the number flux, flow velocity componetns and temperatures. These parameters are given for all points along the radius  $R(I)$  at constant distances  $DR$ . In the low density domain the resolution  $DR$  is much too large compared with the expected mean free path. A different mesh has to be designed for this purpose. It is possible that one cell may be sufficient.
- b. Far Field Condition: The boundary conditions where the gas may be regarded "undisturbed" by the jet are as follows

- Jet gas molecules are allowed to go out the simulation region (these molecules will be regarded as "lost" molecules).
  - Ambient gas is allowed to enter the simulated region according with their thermal velocity and number density.
- c. Solid Wall Bondaries. The solid wall may be assumed to have a constant temperature  $T_w$ . Incident molecules of either species are reflected back from the wall. Different models of collisions with the wall may be employed.
- Elastic collisions 'specular reglection' (this calculation has been included in SIMUL program)
  - Collisions with ideal heat transfer (diffuse reflections)
  - Other models depending on the materials and surface parameters.

The collision with the wall was included as an internal routine in the program SIMUL. If the general molecular simulation contains the program proposed here the collision with the wall will be omitted from SIMUL and become the core of the additional collisionless program. Figure 25 shows the low density (collisionless) region and its boundaries.



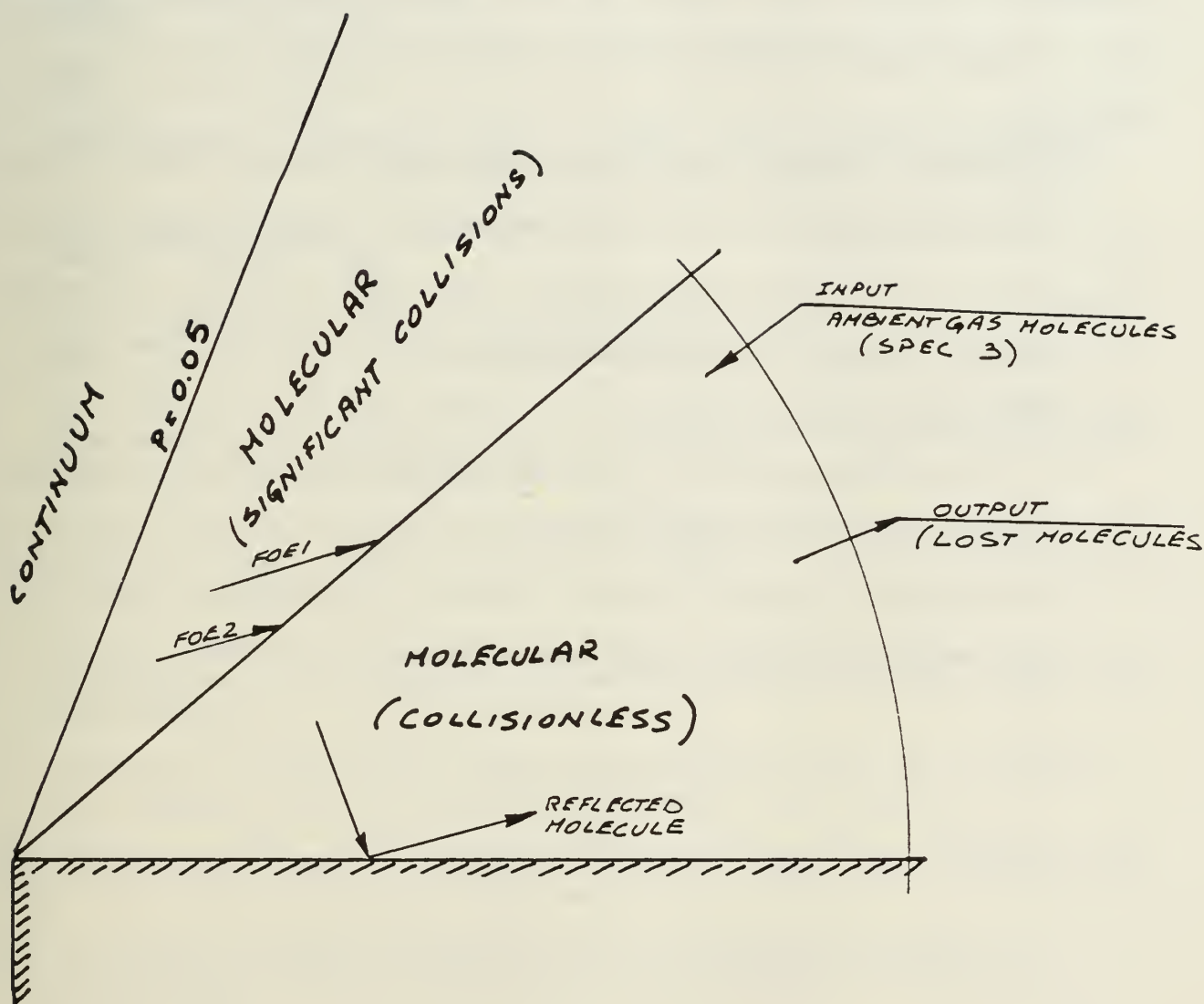


Figure 25. The Low Density Region in the Jet

## SUMMARY OF REPORT

Algorithms for the continuum regime and for the region where molecular collisions are significant have been developed.

Program AXSYM contains the calculation of planar jet flow and axisymmetric ring jet flow. This program supplies data for the limits where the continuum approach become invalid and molecular approach should be employed.

Program SIMUL contains the molecular simulation for the axisymmetric ring flow. This program may run for the whole molecular region to result in the calculation of flux towards the solid wall. For a more efficient simulation it is proposed to design an additional program for the collisionless region where ambient gas may be included.

For the two dimensional flow, program SIMUL may be used after changing the definition of the cell geometry.

To run the whole program it may be required to make separate runs for each sector and store results on the mass storage.

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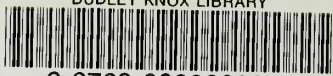
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